Video Article

Immuno-fluorescence Assay of Leptospiral Surface-exposed Proteins

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Abstract

Bacterial surface proteins are involved in direct contact with host cells and in uptake of nutrients from the environment ¹. For this reason, cellular localization can provide insights into the functional role of bacterial proteins. Surface localization of bacterial proteins is a key step towards identification of virulence factors involved in mechanisms of pathogenicity.

Methods for fractionating leptospiral membranes ²⁻⁵ may be selective for a certain class of outer-membrane proteins (OMPs), such as lipoproteins vs. transmembrane OMPs, and therefore lead to misclassification. This likely is due to structural differences and how they are associated to the outer membrane. Lipoproteins are associated with membranes via a hydrophobic interaction between the N-terminal lipid moiety (three fatty acids) and the lipid bilayer phospholipids ^{6, 7}. In contrast, transmembrane OMPs are typically integrated into the lipid bilayer by amphipathic β -sheets arranged in a barrel-like structure ^{8, 9}. In addition, presence of a protein in the outer-membrane does not necessarily guarantee that the protein or its domains are exposed on the surface. Spirochetal outer membranes are known to be fragile and therefore necessitate methods involving gentle manipulation of cells and inclusion of sub-surface protein controls to assess the integrity of the outer membrane.

Here, we present an immunofluorescence assay (IFA) method to directly assess surface exposure of proteins on intact leptospires. This method is based on recognition of leptospiral surface proteins by antigen-specific antibodies. Herein, antibodies specific for OmpL54¹⁰ are detected after binding to native, surface exposed epitopes. Comparison of antibody reactivity to intact versus permeabilized cells enables evaluation of cellular distribution and whether or not a protein is selectively present on leptospiral surface. The integrity of outer membrane should be assessed using antibody to one or more subsurface proteins, preferably located in the periplasm.

The surface IFA method can be used to analyze surface exposure of any leptospiral protein to which specific antibodies are available. Both the usefulness and limitation of the method depends on whether the antibodies employed are able to bind to native epitopes. Since antibodies often are raised against recombinant proteins, epitopes of native, surface-exposed proteins may not be recognized. Nevertheless, the surface IFA method is a valuable tool for studying components of intact bacterial surfaces. This method can be applied not only for leptospires but also other spirochetes and gram-negative bacteria. For stronger conclusions regarding surface-exposure of OMPs, a comprehensive approach involving several cell localization methods is recommended ¹⁰.

Protocol

1. Fixation of *Leptospira* to glass slides

1. *Leptospira interrogans* is a class BSL2 pathogen. Working with live cells it requires appropriate handling, such as wearing gloves, lab-coat and pipetting steps in a sterile-hood.
2. Grow *Leptospira* in EMJH medium¹¹, supplemented with 1% rabbit serum at 30° C until they reach mid- to late-log phase (density of 5×10^7 to 5×10^8 cells/ml) for approximately 6 days.
3. Harvest the culture by centrifugation at $\sim 2000 \times g$ for 7 min at room temperature.
4. Remove the supernatant by aspiration and gently resuspend the pellet in phosphate buffered saline (PBS)-5 mM MgCl₂, pH7.2 to a final concentration of 5×10^8 cells/ml.
5. Add 1 ml of cell suspension to each well of two-well chamber glass slides and incubate at 30° C for 80 min to allow cells to adhere.
6. Carefully remove the liquid containing unbound cells by aspiration.
7. Fix remaining intact bacteria to glass slides by adding 1 ml/well of 2% paraformaldehyde in PBS-5 mM MgCl₂. Incubate for 40 min at 30° C. These slides will be for assessment of surface-exposed proteins.
8. For control slides assessing sub-surface proteins, permeabilize the outer membrane by fixing with 1 ml/well of 100% ice-cold methanol. Incubate at -20° C for 20 min. Methanol acts in several ways: it permeabilizes the outer membrane, denatures the proteins, and fixes cells to glass slides.
9. Remove fixing agents by aspiration.

2. Labeling with specific antibodies

1. Block non-specific binding by adding 1 ml/well of blocking buffer (Difco *Leptospira* Enrichment EMJH). Incubate at 30° C for 90 min.
2. Dilute the specific antibody (immune rabbit sera or mouse monoclonal antibodies, here polyclonal rabbit sera specific for OmpL54¹⁰, FlaA1¹², and OmpL1¹³) and pre-immune rabbit sera or mouse ascetic fluid containing no antibody (when utilized as negative controls) in blocking buffer. Dilutions for each antibody have to be determined empirically depending on antibody titer, antigen-antibody reactivity and abundance of protein in the cell. The usual range is 1:50 to 1:600.

3. Remove the blocking buffer by aspiration and add 1 ml/well of diluted primary antibodies.
4. Incubate at 30° C for 1h.
5. Remove the liquid by aspiration and wash the wells three times with PBS (1 ml/well).

3. Visualization of leptospires

1. Add 1 ml/well of Alexa Fluor 488-labeled secondary antibodies (either goat anti-rabbit IgG or goat anti-mouse IgG) diluted 1:2000 and fluorescent nucleic acid stain, 4'-diamidino-2-phenyl-indole dihydrochloride (DAPI) diluted to a final concentration of 0.25 µg/ml in blocking buffer. This step ensures detection of antibody binding and the presence of all spirochetes independent of antibody binding, respectively.
2. Incubate the slides at 30° C for 45 min.
3. Remove the liquid by aspiration and wash the wells twice with PBS and once with distilled water (1 ml/well).
4. Remove the chambers and the adhesive strip from the glass slides and air-dry for ~10 min.
5. Add ProLong Gold anti-fade mounting medium (2 x 20 µl per slide) and a 24 x 50 mm cover slip.
6. Incubate overnight at room temperature in the dark. This step is necessary to allow the mounting medium to cure (harden).
7. Seal the cover slip with nail polish.
8. Visualize the staining by fluorescence microscopy using a cyan/blue detection filter for DAPI and a green detection filter for Alexa Fluor 488. To ensure that an accurate representation of the samples is made, make sure you evaluate the entire chamber area for each sample.
9. Record the data by imaging a representative field for each sample. When imaging Alexa Fluor 488 fluorescence, use the same exposure time for all samples. Using a consistent exposure time will allow more accurate comparison of results with test antigens versus controls.

4. Representative Results:

Examples of surface IFA results with *Leptospira interrogans* cells using antibodies against surface and sub-surface proteins are shown in Figure 2. A test is considered positive when substantial fluorescence is detected in samples with intact leptospires for a protein of interest that is surface exposed (Fig. 2A). Usually, in a positive test (here, surface-exposed OmpL54), approximately same fluorescence intensity is observed in both intact and permeabilized cells (Fig. 2A). It is essential to include a negative control for outer-membrane integrity using antibodies against a subsurface, preferably periplasmic protein (e.g. leptospiral FlaA1, Fig. 2B). Only when the data show that the outer membrane has remained intact during the experiment, i.e. antibodies against a sub-surface protein do not react to intact cells (no fluorescence or much weaker fluorescence when compared to permeabilized sample as in Fig. 2B), can it be concluded that the protein(s) of interest is surface exposed. Inclusion of experiments with pre-immune sera (in the case of polyclonal antibodies) as an additional negative control is essential for unambiguous data and should not yield substantial fluorescence in intact or permeabilized cells (Fig. 2A, 2C). DAPI stain is necessary to visualize leptospires when negative results (no Alexa Fluor 488 fluorescence) are observed. Counterstaining with propidium iodide is an acceptable alternative to DAPI.

This method is qualitative rather than quantitative as brighter fluorescence may be due to multiple surface exposed antigenic epitopes being recognized in a single antigen relative to other antigens of equal abundance but with fewer surface exposed epitopes. Therefore the fluorescence intensities can only be used to compare reactivity of the same antibodies, such as intact versus permeabilized cells and not fluorescence intensities between different proteins. This distinction is important when an ambiguous result is obtained where significantly less fluorescence is observed in intact cells than in permeabilized cells (Fig. 2C). Such a result indicates that either the protein is in a sub-surface location (with some non-specific background staining of intact cells), the antibodies are binding preferentially to non-native epitopes that are exposed after methanol permeabilization or a protein might have multiple localization sites as described for Erp and Elp proteins of another spirochete, *Borrelia burgdorferi*¹⁴. In either case, an ambiguous IFA result such as this requires alternative approaches such as surface biotinylation or surface proteolysis to determine whether the protein of interest is surface-exposed or not¹⁰.

Figure 1. Flow chart for the surface immuno-fluorescence assay of *Leptospira interrogans* spirochetes. First, 1 ml of cell suspension is added to each well of two-well chamber slides (at least two slides for each experiment) and incubated for leptospires to adhere. Next, leptospires are fixed to chamber slides by adding 1 ml/well of 2% paraformaldehyde (PFA) for samples with intact outer membrane (OM) and 1 ml/well of 100% cold methanol for samples with permeabilized OM. Next, leptospires are incubated with antibodies recognizing surface-exposed proteins along with control antibodies for subsurface proteins (Primary Ab's) at 30° C for 90 min. Then, primary antibodies are detected by incubation with 1 ml/well of Alexa Fluor 488 conjugated secondary antibodies. Finally, cells are visualized by fluorescence microscopy.

Figure 2. Surface immunofluorescence analysis of leptospiral proteins. *Leptospira interrogans* spirochetes were probed with immune and pre-immune sera, and negative results were paired with a DAPI counterstain to demonstrate the presence of spirochetes. A green fluorescence filter was used to detect Alexa Fluor 488 labeled secondary antibody binding to surface exposed proteins and blue/cyan fluorescence filter to detect DAPI binding to cell DNA. A) Leptospires probed with antibodies recognizing a surface exposed protein, OmpL54¹⁰. B) Cells probed with antibodies against a periplasmic subsurface protein, FlaA1 (negative control). C) Cells probed with antibodies against leptospiral porin, OmpL1. Binding of rabbit sera to leptospires were detected with Alexa Fluor 488 conjugated goat anti-rabbit IgG fragments. All images are taken after 4 sec long exposure.

Discussion

Our surface IFA technique is similar to that used to demonstrate the surface exposure of various borrelial^{15, 16} and leptospiral^{12, 17} proteins. The details of the surface IFA method described here are designed to minimize the manipulation of cells in an effort to maintain outer membrane integrity while taking advantage of the tendency of *Leptospira* cells to adhere to glass slides. The method involves a lower concentration (2% versus 4%) of paraformaldehyde, washing with outer-membrane stabilizing buffer (PBS+5mM MgCl₂) and fewer washing steps than previously published protocols^{12, 17}. In addition, our surface IFA method emphasizes the importance of controls that are essential for accurate data interpretation. A limitation of the method is that it requires the availability of specific antibodies that are capable of recognizing surface-exposed epitope(s) of the protein of interest. In some cases, steric hindrance by other surface proteins or lipopolysaccharides can prevent antibody-antigen formation. Nevertheless, the surface IFA method is a relatively straightforward technique for directly assessing surface exposure of proteins and can be used either as a stand alone approach or in concert with other techniques, such as immunogold labeling, surface proteolysis, etc.

Disclosures

No conflicts of interest declared.

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References

1. Achouak, W., Heulin, T. & Pages, J.M. Multiple facets of bacterial porins. *FEMS Microbiol Lett* **199**, 1-7 (2001).
2. Haake, D.A. & Matsunaga, J. Characterization of the leptospiral outer membrane and description of three novel leptospiral membrane proteins. *Infect Immun* **70**, 4936-4945 (2002).
3. Haake, D.A. *et al.* Changes in the surface of *Leptospira interrogans* serovar grippityphosa during *in vitro* cultivation. *Infect Immun* **59**, 1131-1140 (1991).
4. Nally, J.E. *et al.* Purification and proteomic analysis of outer membrane vesicles from a clinical isolate of *Leptospira interrogans* serovar Copenhageni. *Proteomics* **5**, 144-152 (2005).
5. Zuerner, R.L., Knudtson, W., Bolin, C.A. & Trueba, G. Characterization of outer membrane and secreted proteins of *Leptospira interrogans* serovar pomona. *Microb Pathog* **10**, 311-322 (1991).
6. Cullen, P.A., Haake, D.A. & Adler, B. Outer membrane proteins of pathogenic spirochetes. *FEMS Microbiol Rev* **28**, 291-318 (2004).
7. Haake, D.A. Spirochaetal lipoproteins and pathogenesis. *Microbiology* **146** (Pt 7), 1491-1504 (2000).

8. Koebnik, R., Locher, K.P. & Van Gelder, P. Structure and function of bacterial outer membrane proteins: barrels in a nutshell. *Mol Microbiol* **37**, 239-253 (2000).
9. Schulz, G. The structures of general porins. (Benz, R., ed.), *WILEY-VCH Verlag GmbH & Co. KGaA, Weinheim/Germany* (2004).
10. Pinne, M. & Haake, D.A. A comprehensive approach to identification of surface-exposed, outer membrane-spanning proteins of *Leptospira interrogans*. *PLoS One* **4**, e6071 (2009).
11. Johnson, R.C. & Rogers, P. Metabolism of leptospires. II. The action of 8-azaguanine. *Can J Microbiol* **13**, 1621-1629 (1967).
12. Cullen, P.A. *et al.* Surfaceome of *Leptospira* spp. *Infect Immun* **73**, 4853-4863 (2005).
13. Haake, D.A. *et al.* Molecular cloning and sequence analysis of the gene encoding OmpL1, a transmembrane outer membrane protein of pathogenic *Leptospira* spp. *J Bacteriol* **175**, 4225-4234 (1993).
14. Hefty, P.S., Jolliff, S.E., Caimano, M.J., Wikel, S.K. & Akins, D.R. Changes in temporal and spatial patterns of outer surface lipoprotein expression generate population heterogeneity and antigenic diversity in the Lyme disease spirochete, *Borrelia burgdorferi*. *Infect Immun* **70**, 3468-3478 (2002).
15. Noppa, L., Östberg, Y., Lavrinovich, M. & Bergström, S. P13, an integral membrane protein of *Borrelia burgdorferi*, is C-terminally processed and contains surface-exposed domains. *Infect Immun* **69**, 3323-3334 (2001).
16. Parveen, N. & Leong, J.M. Identification of a candidate glycosaminoglycan-binding adhesin of the Lyme disease spirochete *Borrelia burgdorferi*. *Mol Microbiol* **35**, 1220-1234 (2000).
17. Ristow, P. *et al.* The OmpA-like protein Loa22 is essential for leptospiral virulence. *PLoS Pathog* **3**, e97 (2007).