



Extracellular Actin Is a Receptor for *Mycoplasma hyopneumoniae*

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Mycoplasma hyopneumoniae, an agriculturally important porcine pathogen, disrupts the mucociliary escalator causing ciliostasis, loss of ciliary function, and epithelial cell death within the porcine lung. Losses to swine production due to growth rate retardation and reduced feed conversion efficiency are severe, and antibiotics are used heavily to control mycoplasmal pneumonia. Notably, little is known about the repertoire of host receptors that *M. hyopneumoniae* targets to facilitate colonization. Here we show, for the first time, that actin exists extracellularly on porcine epithelial monolayers (PK-15) using surface biotinylation and 3D-Structured Illumination Microscopy (3D-SIM), and that *M. hyopneumoniae* binds to the extracellular β -actin exposed on the surface of these cells. Consistent with this hypothesis we show: (i) monoclonal antibodies that target β -actin significantly block the ability of *M. hyopneumoniae* to adhere and colonize PK-15 cells; (ii) microtiter plate binding assays show that *M. hyopneumoniae* cells bind to monomeric G-actin in a dose dependent manner; (iii) more than 100 *M. hyopneumoniae* proteins were recovered from affinity-chromatography experiments using immobilized actin as bait; and (iv) biotinylated monomeric actin binds directly to *M. hyopneumoniae* proteins in ligand blotting studies. Specifically, we show that the P97 cilium adhesin possesses at least two distinct actin-binding regions, and binds monomeric actin with nanomolar affinity. Taken together, these observations suggest that actin may be an important receptor for *M. hyopneumoniae* within the swine lung and will aid in the future development of intervention strategies against this devastating pathogen. Furthermore, our observations have wider implications for extracellular actin as an important bacterial receptor.

OPEN ACCESS

Edited by:

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Received: 29 November 2017

Accepted: 12 February 2018

Published: 27 February 2018

Citation:

Raymond BBA, Madhkoor R, Schleicher I, Uphoff CC, Turnbull L, Whitchurch CB, Rohde M, Padula MP and Djordjevic SP (2018) Extracellular Actin Is a Receptor for *Mycoplasma hyopneumoniae*. *Front. Cell. Infect. Microbiol.* 8:54. doi: 10.3389/fcimb.2018.00054

Keywords: extracellular actin, mycoplasma infections, host-pathogen interactions, bacterial adhesins, super resolution microscopy

INTRODUCTION

Mycoplasma hyopneumoniae, the etiological agent of porcine enzootic pneumonia (PEP; Mare and Switzer, 1965; Goodwin et al., 1967), remains one of the most economically significant pathogens that causes reduced growth rate along with lower feed conversion efficiency in swine. Attributing losses to pork production incurred by *M. hyopneumoniae* are complicated by polymicrobial infections but recent estimates for US production systems are in the order of \$US10 per

head (Holst et al., 2015). This is significantly more than US estimates of \$200 million to \$1 billion p.a. reported earlier (Clark et al., 1993). On top of this, these estimates fail to account for the environmental impact caused by the release of multiple antibiotic resistant bacterial populations and significant quantities of unmetabolized antimicrobials into effluent ponds, which are in turn used to fertilize agricultural land in many parts of the world, particularly in China; the largest producer and consumer of pork (Wyrsh et al., 2015; Zhu et al., 2015). *M. hyopneumoniae* has an affinity for receptors on the surface of cilia that line the epithelium in the upper respiratory tract of pigs and destroys the mucociliary escalator creating a favorable environment for secondary bacterial infections (Ciprián et al., 1988; Caruso and Ross, 1990; Marois et al., 2007). *M. hyopneumoniae* can also potentiate disease caused by porcine reproductive and respiratory syndrome virus (PRRSV), swine influenza virus (SIV), and porcine circovirus type 2 (PCV2) (Thacker et al., 2000, 2001; Pallarés et al., 2002; Opriessnig et al., 2004). Tetracyclines, macrolides, lincosamides, fluoroquinolones, and aminoglycosides are used widely to treat disease caused by *M. hyopneumoniae* but a more diverse combination of antibiotics is used to prevent polymicrobial respiratory infections in pigs (Stipkovits et al., 2001; Maes et al., 2008). Consequently, disease caused by *M. hyopneumoniae* is one of the major drivers of antibiotic consumption in swine production globally. Thus, there is a pressing need to develop alternatives to antimicrobials to control pathogens that inflict major economic losses in intensively-reared food animals.

Studies over the past 15 years have focussed on understanding how *M. hyopneumoniae* attaches to cilia and colonizes the respiratory epithelium. The P97 and P102 family of multifunctional cilium adhesins are highly expressed (Pendarvis et al., 2014) on the surface of *M. hyopneumoniae* as cleavage fragments that bind several host molecules including highly sulfated glycosaminoglycans (GAGs), fibronectin, and plasminogen (Burnett et al., 2006; Wilton et al., 2009; Deutscher et al., 2010; Seymour et al., 2010, 2011, 2012; Bogema et al., 2011, 2012; Raymond et al., 2013, 2015; Tacchi et al., 2014, 2016). GAGs decorate the surface of cilia within the swine respiratory tract (Erlinger, 1995) and are primary receptors for adhesins on the surface of *M. hyopneumoniae* (Burnett et al., 2006; Jenkins et al., 2006; Wilton et al., 2009; Deutscher et al., 2010; Seymour et al., 2010, 2011, 2012; Bogema et al., 2011, 2012; Raymond et al., 2013, 2015; Tacchi et al., 2014; Jarocki et al., 2015). Upon colonization, *M. hyopneumoniae* induces ciliostasis, loss of cilia, and epithelial cell death (DeBey and Ross, 1994) but it is unknown if *M. hyopneumoniae* can colonize after these events. Furthermore, the identities of other cell surface receptors, especially after ciliary removal, that are targeted by *M. hyopneumoniae* are poorly understood. One such receptor that is of particular interest is the major cytoskeletal protein actin that has been shown to be bound by several bacterial pathogens such as group B streptococcus, *Mycoplasma suis*, and *Legionella pneumophila* (Boone and Tyrrell, 2012; Bugalhão et al., 2015; Zhang et al., 2015). Actin is potentially underappreciated as a bacterial receptor and is reported to be expressed on the surface of a wide range of eukaryote cells (Chen et al., 1978;

Owen et al., 1978; Jones et al., 1979; Sanders and Craig, 1983; Rosenblatt et al., 1985a; Bach et al., 1986; Pardridge et al., 1989; Por et al., 1991; Dudani and Ganz, 1996; Smalheiser, 1996; Miles et al., 2006; Sandiford et al., 2015; Fu et al., 2017; Sudakov et al., 2017). Studies that seek to investigate the ability of *M. hyopneumoniae* to bind to actin have not been undertaken.

Porcine kidney epithelial-like (PK-15) cells were established as a cell line to study the *in vitro* adherence of *M. hyopneumoniae* (Zielinski et al., 1990; Burnett et al., 2006; Wilton et al., 2009; Deutscher et al., 2010; Seymour et al., 2010, 2011). Here, using 3D-structured illumination microscopy (3D-SIM), we show *M. hyopneumoniae* binds to a subset of actin filaments that are expressed on the surface of porcine kidney epithelial cells (PK-15 cells). We assessed the ability of *M. hyopneumoniae* proteins to bind actin using microtiter plate binding assays, affinity chromatography, microscale thermophoresis, immunofluorescence microscopy, and ligand blotting. We also examined the ability of monoclonal antibodies that target β -actin (mAb $_{\beta$ -act) to compete for actin-binding sites on the surface of *M. hyopneumoniae*.

METHODS AND MATERIALS

Bacterial Strains and Culture

Mycoplasma hyopneumoniae strain 232 was grown in modified Friis medium (Scarman et al., 1997) as previously described (Bereiter et al., 1990). Broth cultures were typically incubated at 37°C for ~16 h until the medium turned yellow. For growth on solid agar, cells were plated onto Friis agar as previously described (Kobisch and Friis, 1996).

Protein Expression, Purification, and Generation of Antisera

Purified P97₂₃₂ fragments were expressed and purified as previously described (Raymond et al., 2015). Antiserum against F2_{p94-j} was generated in a previous study and with titers determined empirically (Jenkins et al., 2006).

Quantitative Adherence and Inhibition Assays

Porcine kidney epithelial-like monolayers (PK-15) were grown to semi-confluency and split into 12-well microtiter plates at ~10⁵ cells/ well and left to adhere overnight. For infection, an overnight *M. hyopneumoniae* culture of strain 232 was washed twice in PBS and resuspended in 25 mM HEPES in DMEM containing 5% fetal bovine serum (infection medium) and incubated at 37°C for 2 h. A proportion of these cells were diluted in fresh Friis broth and plated onto Friis agar to establish cell numbers prior to infection. *M. hyopneumoniae* cells were then added to each well containing PK-15 cells at a multiplicity of infection (MOI) of 0.85. *M. hyopneumoniae* cells were allowed to adhere for 16 h at 37°C/ 5% CO₂. Non-adherent cells were removed by washing in PBS followed by incubating the wells in TrypLE™ (Thermo Fisher Scientific) for 20 min at 37°C to remove adherent PK-15 and *M. hyopneumoniae* cells. Cells were then removed by gently pipetting to create a relatively homogenous suspension followed by dilution in

Friis broth to 10^{-1} and 10^{-2} and plating onto Friis agar. Plates were incubated for a minimum of 7 days at 37°C / 5% CO_2 prior to counting under a stereomicroscope. Inhibition experiments were performed by pre-incubating PK-15 cells with a 1:100 dilution of monoclonal antibodies against β -actin (IgG2A mAb $_{\beta\text{-act}}$; $\sim 5\text{--}20\ \mu\text{g ml}^{-1}$ mAb, Sigma-Aldrich) for 2 h prior to the addition of *M. hyopneumoniae* cells. To ensure that IgG2A itself has no effect on the adherence of *M. hyopneumoniae* cells to PK-15 monolayers, isotype control experiments were performed. PK-15 monolayers, seeded onto coverslips as described below, were pre-incubated with either mAb $_{\beta\text{-act}}$, murine IgG2A (Sigma-Aldrich), or no antibody for 2 h prior to infection with *M. hyopneumoniae*. Samples were then processed for immunofluorescence microscopy as described below. Slides were imaged using an Olympus BX51 Upright Epi Fluorescence Microscope at $20\times$ magnification and a constant 100 ms exposure, using the GFP filter to capture adherent *M. hyopneumoniae* cells labeled with CFTM 488. Five fields of view were captured per sample, equating to the adherence of *M. hyopneumoniae* cells across approximately 1,500 PK-15 cells per field of view. Image thresholding was performed using ImageJ and the mean fluorescence was calculated on the binary images generated. GraphPad Prism 7 was used to plot the data, including the standard error of the mean, and to perform the statistical analyses (unpaired *t*-test).

Immunofluorescence Microscopy of PK-15 Monolayers Infected with *M. hyopneumoniae* Cells

Experiments were performed as described above except that PK-15 cells were split into microtiter plates containing glass coverslips (15 mm, 170 μm thickness; ProSciTech). Following infection, samples were washed $3\times$ in PBS and fixed in 2% methanol-free paraformaldehyde overnight at 4°C . Prior to commencing the following steps, samples were washed $3\times$ in PBS and all antibody dilutions were made in 1% BSA in PBS. Excess aldehydes were quenched in 100 mM glycine in PBS for 5 min at RT. Non-specific binding sites were blocked in 2% BSA/PBS for 1 h at RT. *M. hyopneumoniae* cells were labeled using polyclonal F2_{P94-J} rabbit antisera that recognizes the C-terminus of P97, incubated at a dilution of 1:500 for 1 h at RT. A 1:1,000 dilution of anti-rabbit CFTM 488- or 568-labeled secondary antibody (Biotium) was incubated for 1 h at RT. Cells were permeabilized in 0.5% (v/v) Triton X-100 in PBS for 5 min. DAPI (Roche) was then added for 5 min at RT to stain nucleic acids. Phalloidin conjugated to CFTM 568 (Biotium) was then added for 30 min at RT to stain filamentous (F-actin) actin. Coverslips were then mounted onto glass microscope slides in VECTASHIELD[®] (Vector Laboratories). Samples were imaged using a Nikon A1 Confocal Laser Scanning Microscope and a V3 DeltaVision OMX 3D-Structured Illumination Microscopy Imaging System (Applied Precision, GE Healthcare) as previously described (Strauss et al., 2012). Images were processed using Imaris Scientific 3D/4D image processing software (Bitplane AG, Zurich, Switzerland).

Immunofluorescence Microscopy of Surface Exposed Actin in PK-15 Monolayers

Experiments were performed identically to those described above except that a portion of PK-15 cells were not infected with *M. hyopneumoniae*. Surface actin was labeled prior to permeabilization using a murine monoclonal antibody against β -actin (mAb $_{\beta\text{-act}}$, IgG2a isotype) at a dilution of 1:100 for 1 h at RT, followed by incubation with a 1:1,000 dilution of anti-murine CFTM 488-conjugated secondary antibodies (Biotium) for 1 h at RT. A number of control experiments were performed in order to confirm the specificity of mAb $_{\beta\text{-act}}$. For surface exposed actin and to rule out the possibility of non-specific binding. Initially, an isotype control was performed using murine IgG2a that was diluted 1:100 in 2% BSA/PBS and incubated for 1 h at RT, followed by incubation with a 1:1,000 dilution of anti-murine CFTM 488-conjugated secondary antibodies (Biotium) for 1 h at RT. Secondary antibody controls were performed as described above, without primary antibody incubation. Additional controls were performed using mitochondria as an intracellular marker of permeabilization/compromization of the membrane. Uninfected PK-15 cells were fixed in paraformaldehyde as described above and were then either permeabilized in 0.5% Triton X-100 for 5 min or left unpermeabilized. Murine monoclonal antibodies against mitochondria (Abcam) were diluted 1:100 in 2% BSA/PBS and incubated overnight at 4°C , followed by incubation with a 1:1,000 dilution of anti-murine CFTM 488-conjugated secondary antibodies (Biotium) for 1 h at RT. Prior to mounting, all samples were stained with DAPI and imaged using an Olympus BX51 Upright Epi Fluorescence Microscope.

Infection Experiments for Scanning Electron Microscopy

PK-15 cells were maintained in Dulbecco's modified eagle medium (DMEM, PAA Laboratories, Pasching, Austria) containing 10% fetal calf serum, 2 mM L-glutamine, 100 U ml^{-1} penicillin, and 0.1 mg ml^{-1} streptomycin in culture dishes (9 cm in diameter, TPP, Trasadingen, Switzerland) at 37°C in 5% CO_2 . Cells were harvested by trypsinization, resuspended in DMEM with 5% FCS and 2 mM L-glutamine and seeded into 24-well plates containing sterile 12 mm glass cover slips at 1×10^5 cells/well. After growing overnight the medium was changed to infection medium (DMEM with 25 mM HEPES, 2 mM L-glutamine, 5% FCS) by adding 500 μl medium per well. 6 ml cultures of *M. hyopneumoniae* were harvested by centrifugation (14,000 rpm for 5 min, Eppendorf, Hamburg) and resuspended in 1 ml fresh infection medium for 2 h at 37°C on a roller, centrifuged again and resuspended in 120 μl infection medium. Ten to thirty microliters of the bacterial suspension were added per well and incubated at 37° in 5% CO_2 for 4–48 h for infection. After different infection times samples were washed once in infection medium in a beaker and subsequently fixed with 2% glutaraldehyde and 5% formaldehyde in cacodylate buffer (100 mM sodium dimethylarsenic acid, 10 mM MgCl_2 , 10 mM CaCl_2 , 90 mM sucrose, pH 6.9) for 1 h on ice. Samples were then washed with TE buffer (20 mM TRIS, 2 mM EDTA,

pH 7.0), dehydrated in a graded acetone series (10, 30, 50, 70, 90, 100%) for 10 min each step on ice and critical-point dried using liquid CO₂. Samples were made conductive by sputter coating with a thin gold film. Samples were examined in a Zeiss field emission scanning electron microscope DSM982 Gemini at an acceleration voltage of 5 kV using the Everhart-Thornely SE-detector and the inlens SE-detector in a 50:50 ratio. Images were recorded on a MO-disk.

Avidin Purification of Interacting Proteins

The purification of porcine proteins from PK-15 monolayers that bind to proteins expressed on the surface of *M. hyopneumoniae* proteins was performed using methods described previously (Raymond et al., 2013). *M. hyopneumoniae* cells from a 250 ml culture were pelleted, washed, and resuspended in PBS containing 2 mM EZ-link Sulfo-NHS-Biotin for 30 s on ice. This ensured that *M. hyopneumoniae* cells did not lyse, and any remaining biotin molecules were quenched in 100 mM glycine for 10 min at RT. Cells were washed twice by centrifugation in PBS to remove excess glycine. Cells were gently lysed in 0.5% Triton X-100 in Tris-HCl pH 7.6, 150 mM NaCl with protease inhibitors on ice with vortexing and water bath sonication. Insoluble material was removed by centrifugation at 16,000 × *g* for 10 min. The cleared lysate was incubated with avidin agarose (Thermo Scientific) for 1 h at RT on a rotating wheel. The mixture was packed into a column and any unbound *M. hyopneumoniae* proteins were removed by thoroughly washing with PBS.

PK-15 cells were grown and harvested as previously described, without biotinylation (Raymond et al., 2013). The cleared protein lysate was incubated with the biotinylated-*M. hyopneumoniae*-avidin mixture overnight on a rotating wheel at 4°C. The mixture was packed into a column and the procedure was performed as previously described (Raymond et al., 2013).

Biotinylation of PK-15 Surface Proteins

The biotinylation of PK-15 cells was performed as previously described with slight modifications (Raymond et al., 2013). PK-15 cells were grown to semi-confluency in a 175 cm² flask, washed in PBS, and incubated with PBS containing 2 mM EZ-link Sulfo-NHS-Biotin for 30 s on ice. Excess biotin molecules were quenched in 100 mM glycine for 10 min at RT and cells were washed twice in PBS to remove excess glycine. Cells were gently lysed in 0.5% Triton X-100 in Tris-HCl pH 7.6, 150 mM NaCl with protease inhibitors on ice with vortexing and water bath sonication. Insoluble material was removed by centrifugation at 16,000 × *g* for 10 min. The cleared lysate was incubated with avidin agarose (Thermo Scientific) for 1 h at RT on a rotating wheel. The mixture was packed into a column and the washes, elutions, and preparations and identifications using LC-MS/MS were performed as previously described (Raymond et al., 2013).

Preparation of Monomerized Actin

For binding experiments we utilized actin from bovine muscle (Sigma-Aldrich), likely α- or γ-actin. Sequence alignments performed on porcine α-, β-, and γ- isoforms demonstrated that they share 92.5% sequence identity (Supplementary Figure 5) and only differ substantially in the N-terminal 20 amino acids.

This suggests that for the purposes of binding experiments, the utilization of actin from muscle (presumably α- and γ-actin) is sufficient to mimic β-actin binding. Additionally, bovine and porcine actin share 100% sequence identity, making it an appropriate substitution for investigating binding.

Actin was solubilized in 8 M Urea/ 20 mM Triethylammonium bicarbonate pH 8.0. Actin polymerization is dependent on disulfide bridges (Otterbein et al., 2001) and so in order to maintain actin in its monomerized form (G-actin), cysteine residues were reduced and alkylated in 20 mM acrylamide monomers and 5 mM Tributylphosphine for 90 min. Actin monomers were labeled in 20 × molar excess Sulfo-NHS-LC-Biotin for 3 h at RT. Removal of excess biotin and buffer exchange into PBS was performed in a PD-10 Desalting Column (GE Healthcare, Life Sciences) as per manufacturer instructions.

Avidin Purification of Actin-Binding Proteins

Biotinylated actin was incubated with avidin agarose (Thermo Scientific) on a rotating wheel for 5 h. The slurry was packed into a column and the flow through collected. Unbound actin was thoroughly washed with PBS.

Native *M. hyopneumoniae* whole cell lysates were prepared as previously described (Raymond et al., 2013). The native lysate was incubated with the biotinylated actin-avidin agarose mixture overnight on a rotating wheel at 4°C. The mixture was packed into a column and the flow through collected. Unbound proteins were thoroughly washed and collected in PBS and interacting proteins were collected in 30% acetonitrile/ 0.4% trifluoroacetic acid. The elutions were concentrated in a 3,000 Da cutoff filter and acetone precipitated at −20°C for 30 min. Proteins were pelleted by centrifugation at 25,000 × *g* at 4°C for 30 min. Protein was resuspended in SDS sample buffer and separated by 1D SDS-PAGE. Proteins were in-gel trypsin digested and analyzed by LC-MS/MS.

Whole Cell ELISA

A 48 h Mhp232 culture was washed × 3 in PBS and fixed in 1% paraformaldehyde for 1 h at room temperature. Fixed cells were diluted in carbonate-coating buffer (18 mM NaHCO₃, 27 mM Na₂CO₃; pH 9.5) and an equivalent of 100 μL of the original culture was added to each well of a MaxiSorp™ 96-well plate (Nunc®) and centrifuged at 2,000 rpm for 15 min. 20 mg ml^{−1} of BSA was used as a negative control for this experiment. Wells were washed × 3 in wash solution (0.2% tween-20 in PBS). Wells were blocked in 2% BSA in PBS for 1 h at room temperature. Wells were washed × 3 and incubated with 0, 1, 5, 10, 20, and 50 μg ml^{−1} of biotinylated bovine actin (Sigma-Aldrich) in 1% BSA in PBS for 1.5 h at room temperature. Wells were washed × 10 and incubated with a 1:2,000 dilution of ExtrAvidin®-Peroxidase (Sigma-Aldrich) in 1% BSA in PBS for 1 h at room temperature. Wells were washed × 10 and developed using TMB Substrate Reagent Set (BD Biosciences OptEIA™) and read at 650 nm after 15 and 30 min on a PowerWave HT Microplate Spectrophotometer (Bio-Tek). Background binding observed in wells without cells were subtracted from all readings. Data was

analyzed using GraphPad Prism 6 software and has been plotted as the average of triplicate experiments.

Microscale Thermophoresis

Actin monomers were fluorescently labeled with amine reactive NT-647 dye (NanoTemper) in a three-fold molar excess for 30 min and excess dye was removed using a desalting column (NanoTemper). Tween-20 (0.5%) was added to prevent sample aggregation. Non-fluorescent P97 recombinant fragments F1_{P97–232}, F3_{P97–232}, and F4_{P97–232} were titrated against 100 nM fluorescent actin in a 1:1 serial dilution starting from 30 μ M for F3_{P97–232} and F4_{P97–232} and 11 μ M for F1_{P97–232} and samples were loaded into Monolith NT.115™ (NanoTemper) hydrophilic capillaries. Capillary scans were performed to confirm that no sample aggregation was occurring and that the fluorescence intensity was within the range of 80–1,500 counts. The data presented here was generated at 40% MST laser power measuring the thermophoretic movement after the MST laser is switched off for 30 s. NanoTemper analysis software was used for analysis and to generate K_D values.

One- and Two-Dimensional Gel Electrophoresis and Ligand Blotting

1D and 2D SDS-PAGE was performed as previously described (Raymond et al., 2013) with minor modifications. Once *M. hyopneumoniae* proteins were transferred to PVDF membrane and blocked, biotinylated actin, or biotinylated PK-15 lysate prepared as previously described (Raymond et al., 2013) was incubated with the membrane overnight at 4°C. Membranes were probed with a 1:2,000 dilution of ExtrAvidin (Sigma-Aldrich) and developed with diaminobenzidine (Sigma-Aldrich).

1D LC-MS/MS Using QTOF

Methods were performed as described previously with no modifications (Raymond et al., 2013).

MS/MS Data Analysis

MS/MS data files were searched using Mascot as previously described (Raymond et al., 2013) with slight modifications. Data files from avidin purification of interacting proteins were searched for both bacterial and mammalian proteins in order to identify the biotinylated “bait” proteins eluted in their respective experiments.

RESULTS

M. hyopneumoniae Co-localizes with Actin When Adhering to PK-15 Cells

The pattern of adherence of *M. hyopneumoniae* on PK-15 cell monolayers has not previously been studied by fluorescence microscopy. By using the filamentous actin (F-actin) dye phalloidin to stain the cytoskeleton of PK-15 cells and F2_{P94–J} antibodies that recognize the P97 adhesin, we repeatedly observed *M. hyopneumoniae* cells attaching to PK-15 cells along the leading edges as well as in discrete pockets on the surface that stained intensely with phalloidin

(Figure 1 and Supplementary Figure 1). *M. hyopneumoniae* cells also co-localize with phalloidin-staining filaments that resemble filopodia (Figure 1C). Scanning Electron Microscopy images demonstrated a similar pattern of adherence, with *M. hyopneumoniae* cells preferentially colonizing the leading edges of the monolayer (Figures 2A,B). These observations suggest that *M. hyopneumoniae* cells can adhere to areas that stain with phalloidin; inferring that actin may be a potential receptor. 3D-SIM images of *M. hyopneumoniae* adhering to PK-15 cell monolayers once again demonstrated that regions staining intensely with phalloidin are favored (Figure 2C). A cross-sectional view of the image depicted in Figure 2C shows *M. hyopneumoniae* cells in close association with F-actin (Figure 2D). Given the intimate association between *M. hyopneumoniae* cells and actin, we hypothesized that *M. hyopneumoniae* cells may be targeting extracellular actin during adherence. However, PK-15 cell membranes must first be permeabilized to stain with phalloidin and it remains a challenge to determine if the F-actin making intimate contact with *M. hyopneumoniae* is extracellular.

M. hyopneumoniae Co-localizes with Extracellular β -Actin

There are three main isoforms of actin; α -, β -, and γ -. α - and γ -actins are predominantly found in muscle while β -actin is the most predominant form found in non-muscle cells (Herman, 1993). Each of the actin isoforms exist either as a globular form (monomer), referred to as G-actin, or a polymerized filamentous form known as F-actin. Since our data suggests that *M. hyopneumoniae* co-localizes with F-actin (polymerized form of β -actin) we sought to investigate the localization of actin on the surface of uninfected PK-15 cells using 3D-SIM. To ensure that PK-15 cell monolayers did not become permeabilized during fixation, we labeled paraformaldehyde-fixed PK-15 cells that were permeabilized with Triton X-100 with a monoclonal antibody that recognizes mitochondria and compared the images with unpermeabilized cell preparations (no Triton X-100). Mitochondria were intensively stained in PK-15 cell preparations that were permeabilized using Triton X-100 (Supplementary Figure 2) but remained unlabeled (using identical exposure times) in cell preparations that were not treated with Triton X-100 (Supplementary Figure 2). These experiments indicated that the fixation protocol did not permeabilize PK-15 cell membranes. Additionally, isotype and secondary antibody controls confirmed that non-specific binding interactions were undetectable (Supplementary Figure 2).

3D-SIM images of PK-15 cells that were labeled prior to permeabilization with mAb $_{\beta}$ -act depict extracellular β -actin colocalizing with phalloidin-stained regions of the cell, including defined areas at the leading edge of PK-15 cells (Figures 3A–C). We also observed a pattern of staining where *M. hyopneumoniae* did not colocalize with phalloidin-staining filaments. In methodologically unrelated experiments, we identified tryptic peptides that mapped to β -actin from several discrete bands from an SDS-PAGE gel with molecular masses ranging from 40 kDa (predicted monomeric size of actin) to

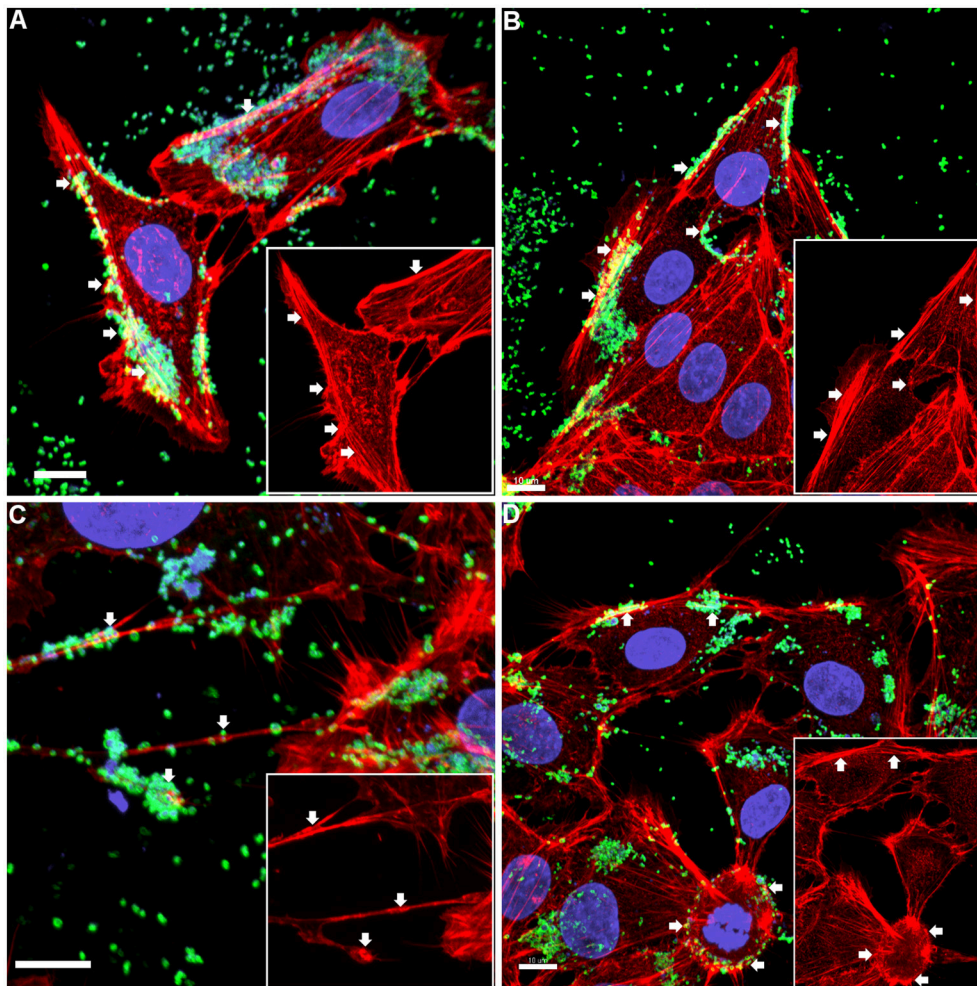


FIGURE 1 | Confocal micrograph overlays of adherence patterns of *M. hyopneumoniae* cells to PK-15 monolayers. *M. hyopneumoniae* cells were labeled with F2p94–J antisera conjugated to CFTM 488 (green), nucleic acids were stained with DAPI (blue), and F-actin was stained using phalloidin (red). *M. hyopneumoniae* cells can be seen adhering to the edges of the monolayer and to cellular projections that appear to be filopodia (white arrows). Inserts in each panel depict the phalloidin channel of each image (more specifically, the areas highlighted in the overlays), highlighting areas that stain intensely for F-actin (white arrows). (A–D) represents images taken from individual replicate experiments. Scale bars are 10 μm .

>100 kDa (polymerized forms of actin). The protein bands were recovered by avidin chromatography from a lysate of PK-15 cells whose surface proteins had been labeled with biotin (Supplementary Figure 3) indicating that extracellular actin exists in multiple forms on the surface of PK-15 cells. When we examined PK-15 cells that had been infected with *M. hyopneumoniae* under the same conditions, we observed the mycoplasma cells colocalizing with extracellular β -actin that was labeled with mAb $_{\beta\text{-act}}$ (Figure 3D). Additionally, we could demonstrate mAb $_{\beta\text{-act}}$ to compete with *M. hyopneumoniae* for surface accessible actin sites, reducing adherence by $\sim 90\%$ compared with non-treated controls (Figure 4B). Isotype control experiments demonstrated that pre-incubating PK-15 monolayers with murine IgG2A prior to infection with *M. hyopneumoniae* had no inhibitory effect on adherence (Supplementary Figure 4).

***M. hyopneumoniae* Surface Adhesins Bind to Extracellular Actin**

To assess the ability of *M. hyopneumoniae* cells to bind to G-actin, we used a microtiter plate binding assay. G-actin bound to the surface of *M. hyopneumoniae* in a dose-dependent manner and reached saturation at $20 \mu\text{g ml}^{-1}$ (Figure 4A).

As part of a larger investigation to determine the identities of epithelial cell receptors that bind to proteins on the surface of *M. hyopneumoniae*, we coupled avidin agarose with surface-accessible *M. hyopneumoniae* membrane proteins that were labeled with sulfo-NHS-LC-biotin and exposed the derivatized agarose to a native PK-15 cell lysate. After extensive washing, the eluents were digested with trypsin and tryptic peptides were sequenced by LC-MS/MS. Actin, and other proteins with cytoskeletal functions such as vimentin, keratin, tubulin, myosin, and tropomyosin (Supplementary Table 1) comprised 13% of the

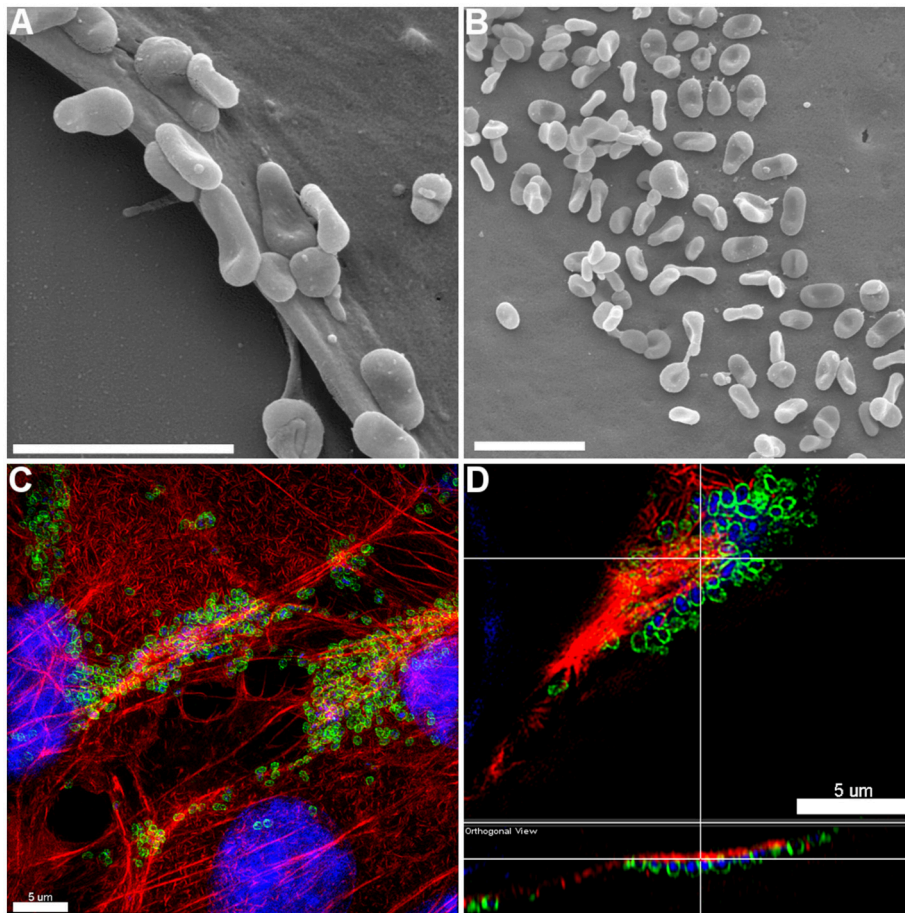


FIGURE 2 | Adherence patterns of *M. hyopneumoniae* cells to PK-15 monolayers. **(A,B)** Scanning electron microscopy (SEM) image of *M. hyopneumoniae* cells adhering to PK-15 monolayers. *M. hyopneumoniae* cells favor the leading edge of epithelial cells **(A)** and select pockets on the top of the cells **(B)**, a pattern consistent with selective localization of epithelial receptors that are a target of adhesins on the surface of *M. hyopneumoniae*. Scale bars in **(A,B)** are 2 μm . **(C,D)** 3D-SIM images of *M. hyopneumoniae* adhering to PK-15 cell monolayers. *M. hyopneumoniae* cells were labeled with F2_{P94-J} antisera conjugated to CFTM 488 (green), nucleic acids were stained with DAPI (blue), and F-actin was stained using phalloidin (red). *M. hyopneumoniae* cells often localize to regions enriched in F-actin. **(D)** Orthogonal view showing *M. hyopneumoniae* cells associating closely with F-actin. Upper panel shows xy orientation and bottom panel shows xz orientation. White lines indicate the same area within the image.

105 PK-15 cell-derived proteins that were identified. To further investigate the role of actin as a potential epithelial cell receptor for *M. hyopneumoniae*, we immobilized biotinylated G-actin on avidin agarose beads and incubated them with a whole cell lysate of *M. hyopneumoniae*. After extensive washing to remove unbound proteins, the column was washed with 2M NaCl (13 times physiological concentration) and the remaining bound proteins were eluted using 0.4% trifluoroacetic acid (**Figure 5D**). 143 *M. hyopneumoniae* proteins were identified by LC-MS/MS (Supplementary Table 2), including most members of the P97 and P102 adhesin families and several cleavage fragments of the archetype cilium adhesin P97. Consistent with data from affinity chromatography experiments, ligand blotting studies showed that biotinylated G-actin bound to many *M. hyopneumoniae* proteins (**Figure 5E**). Of these 143 proteins, 75 (52%) were found in our surfaceome studies of *M. hyopneumoniae* (**Figure 4C** and Supplementary Table 2; Berry et al., 2017). On a cautionary note,

due to the native conditions under which these experiments were performed, some of the 143 proteins may exist as a complex with *bona fide* actin-binding proteins. As such, these proteins should be considered as putative actin-binding proteins. In order to validate the proteins captured by affinity chromatography as *bona fide* actin-binding proteins, we tested several previously described (Raymond et al., 2015) recombinant fragments of P97 for their ability to bind actin using microscale thermophoresis. The recombinant fragments F1_{P97-232} (amino acids 106–758) and F3_{P97-232} (amino acids 768–938) bound actin with K_D 's of 34.2 nM and 1.79 μM , respectively (**Figures 5A,B**). Notably, F4_{P97-232} which contains a multifunctional binding motif with sequence ¹⁰⁷⁰K-K-S-S-L-K-V-K-I-T-V-K¹⁰⁸¹ did not bind actin (**Figure 5C**; Raymond et al., 2015). Taken together, these data indicate that globular β -actin can act as a primary receptor for *M. hyopneumoniae* cells and that a number of surface exposed adhesins are involved in binding to extracellular actin.

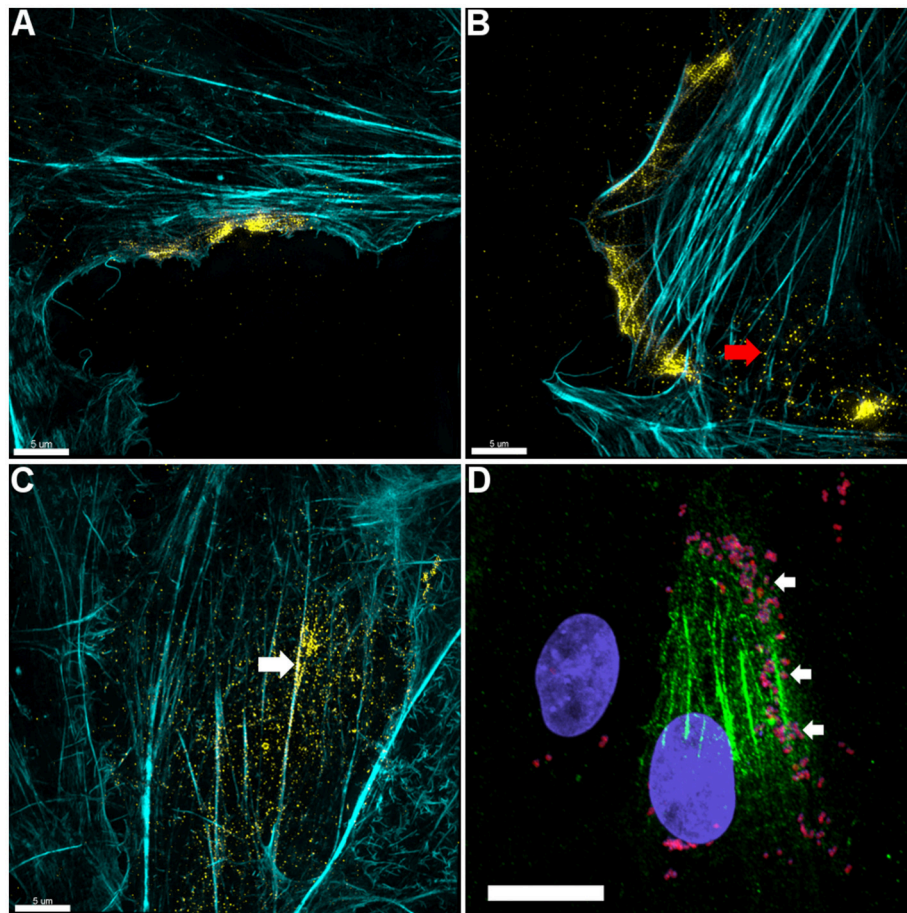


FIGURE 3 | Actin is expressed on the surface of PK-15 cell monolayers. **(A–C)** Localization of actin on the surface of PK-15 cells. Uninfected monolayers were grown to semi-confluency, fixed in methanol-free paraformaldehyde, incubated with mAb $_{\beta}$ -act prior to permeabilization, and conjugated to anti-murine CFTM 488 (yellow). F-actin was then stained using phalloidin (cyan) after permeabilization. Surface exposed actin can be seen accumulating at the leading edge **(A,B)** of cells in the monolayer as well as on the surface of cells along filaments that stained intensely with phalloidin **(C)**. Red arrow in **(B)** indicates what appears to be globular extracellular actin while the white arrow in **(C)** demonstrates extracellular actin that appears to colocalize with F-actin. **(D)** *M. hyopneumoniae* cells adhere on the surface of PK-15 cells in regions enriched with surface actin. *M. hyopneumoniae* cells (magenta colored cells indicated by white arrows) were labeled with F2_{P94–J} antisera conjugated to CFTM 568 (red) and in this case, surface actin appears green. The nuclei of *M. hyopneumoniae* and PK-15 cells were stained with DAPI (blue). Scale bars in **(A–C)** are 5 and 20 μ m in **(D)**.

DISCUSSION

LC-MS/MS analysis of affinity-purified, biotinylated surface proteins derived from PK-15 cells identified tryptic peptides that mapped to actin (Supplementary Figure 3). This observation provided the first clue that actin resides on the surface of PK-15 cells. 3D-SIM of non-permeabilized PK-15 cells that were labeled with mAb $_{\beta}$ -act confirmed the extracellular localization of β -actin and, for the first time, *M. hyopneumoniae* interacting intimately with it (**Figure 1**). Actin localizes on the surface of PK-15 cells in distinct locations, including the cell's leading edge and along actin filaments that stain with phalloidin, a pattern that mirrors the adherence of *M. hyopneumoniae* to PK-15 cells (**Figure 1**). This is the first observation of *M. hyopneumoniae* colocalizing with extracellular actin and implicates it as a putative bacterial receptor on epithelial cell surfaces.

From this study, there are now several lines of evidence that are consistent with the hypothesis that extracellular actin is an important receptor for *M. hyopneumoniae*: (i) mAb $_{\beta}$ -act can inhibit the ability of *M. hyopneumoniae* to adhere to and colonize PK-15 cells by $\sim 90\%$; (ii) microtiter plate binding assays show that *M. hyopneumoniae* cells bind to G-actin in a dose dependent and saturable manner; (iii) ligand blotting studies show biotinylated G-actin binds directly to *M. hyopneumoniae* proteins; (iv) microscale thermophoresis showed that recombinant N-terminal F1_{P97–232} and central fragment F3_{P97–232} of the cilium adhesin P97 bind G-actin with K_D 's of 34.2 nM and 1.79 μ M, respectively; (v) affinity chromatography studies using biotinylated *M. hyopneumoniae* surface proteins as bait identified 105 PK-15 cell proteins including actin and other cytoskeletal proteins such as vimentin, keratin, tubulin, myosin, and tropomyosin; (vi) biotinylated

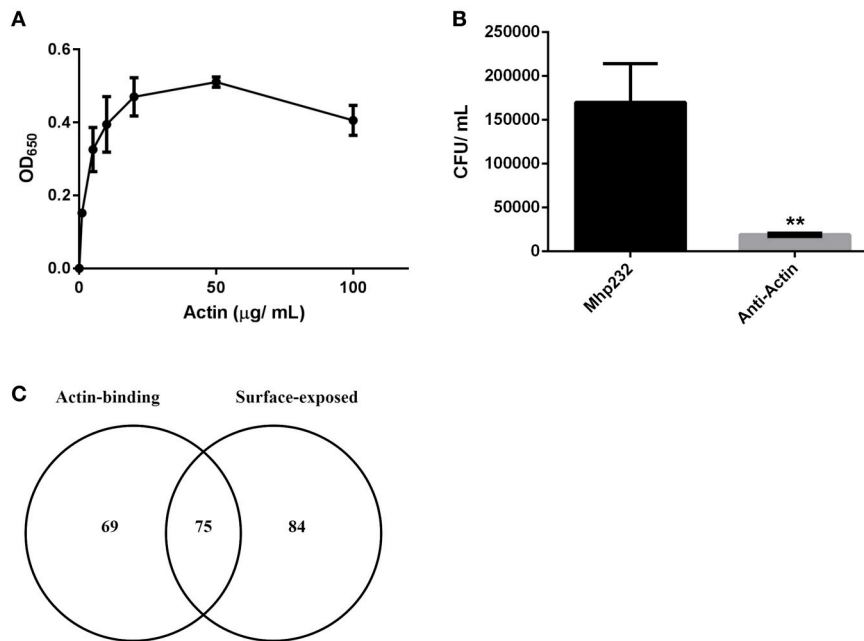


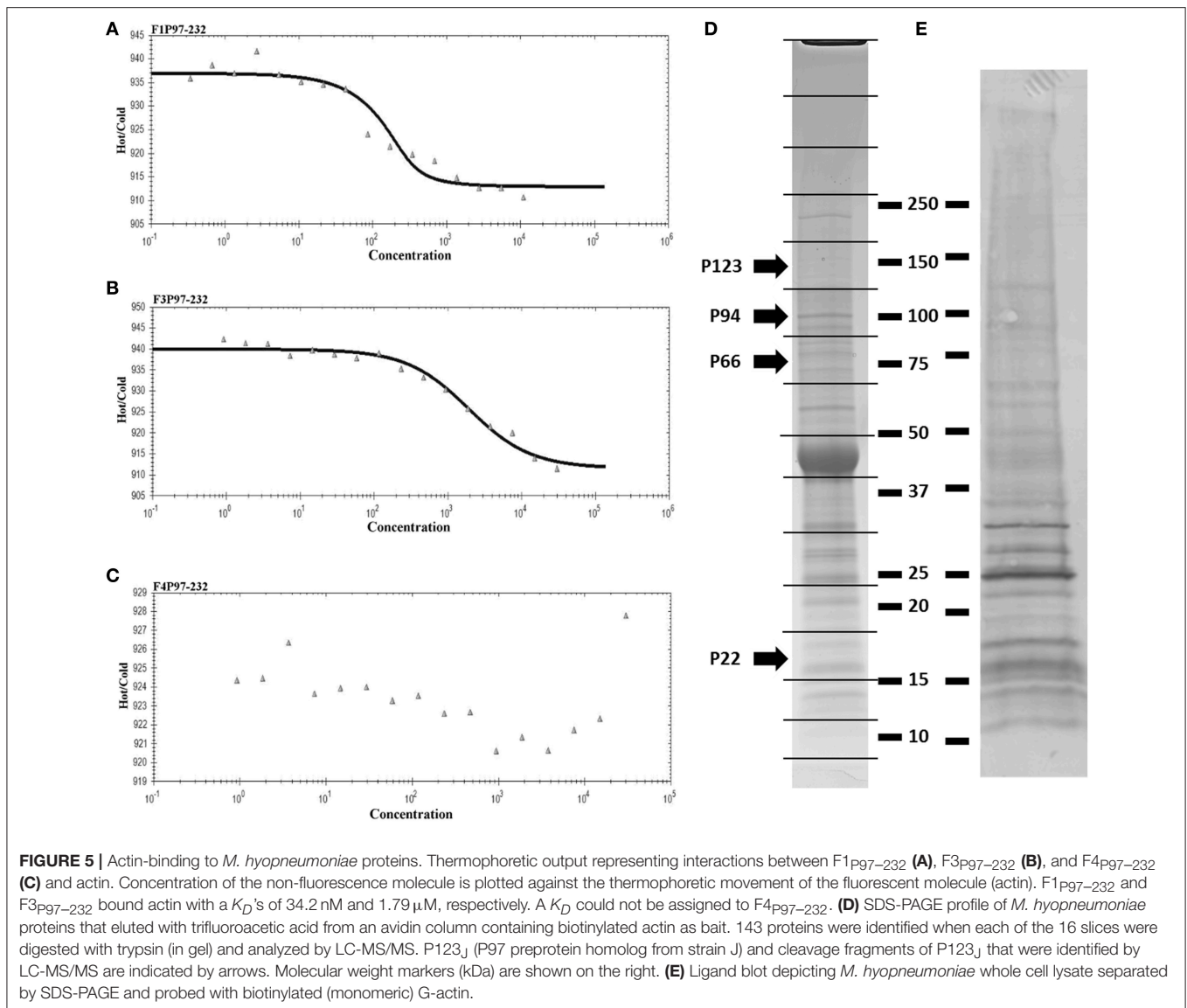
FIGURE 4 | *M. hyopneumoniae* binds actin. **(A)** Microtiter plate binding assay of immobilized *M. hyopneumoniae* cells binding to actin in a dose-dependent and saturable manner. Each data point represents the average of individual triplicate experiments with the standard error of the mean shown. **(B)** Treatment of PK-15 cell monolayers with mAbs that target actin inhibited adherence of *M. hyopneumoniae* by ~90%. The data is presented as the average of triplicate experiments and the standard error of the mean. An unpaired *t*-test was performed with a $p < 0.01$, indicated as **. **(C)** Venn diagram showing the overlap between putative actin-binding proteins purified using actin affinity chromatography, and *M. hyopneumoniae* surface proteins.

G-actin used as bait, recovered 143 *M. hyopneumoniae* proteins including major cilium and extracellular matrix binding proteins. P97 has previously been shown to bind glycosaminoglycans (GAGs), fibronectin, and plasminogen (Minion et al., 2000; Djordjevic et al., 2004; Jenkins et al., 2006; Wilton et al., 2009; Raymond et al., 2015) and now actin. The diversity of molecules presented by epithelial cells that are known to be targeted by adhesins like P97 is consistent with the organism having the capability to adhere to multiple tissue sites and regulate tissue tropism and reinforces the notion that *M. hyopneumoniae* is able to colonize extrapulmonary sites (Lloyd and Etheridge, 1981; Williams and Gallagher, 1981; Yagihashi et al., 1984; Le Carrou et al., 2006; Marois et al., 2007; Woolley et al., 2012). To the best of our knowledge, this is the first study to comprehensively investigate the adherence of a bacterial pathogen to actin; an important cytoskeletal protein that exists in essentially all eukaryote cells.

Porcine epithelial receptors that are targeted by adhesins on the surface of *M. hyopneumoniae* have largely remained uncharacterized. GAGs expressed on the surface of porcine cilia (Erlinger, 1995) have long been known to act as primary receptors for *M. hyopneumoniae* cells (Zhang et al., 1994) and that pre-incubation of *M. hyopneumoniae* cells with cilia or the negatively charged GAGs that decorate cilia can significantly inhibit cilia adherence (Zhang et al., 1994). *M. hyopneumoniae* has also been shown to bind and colocalize with fibronectin at the site of infection (Burnett et al., 2006; Jenkins et al., 2006; Wilton

et al., 2009; Deutscher et al., 2010; Seymour et al., 2010, 2011, 2012; Bogema et al., 2011, 2012; Raymond et al., 2013, 2015; Tacchi et al., 2014, 2016; Jarocki et al., 2015) however, little is known about receptors that may be conducive to facilitating chronic infection such as those that are expressed on the surface of denuded epithelium. Coupling biotinylated surface proteins from *M. hyopneumoniae* onto avidin agarose enabled the recovery of PK-15 receptors that were identified using LC-MS/MS. 105 proteins were identified that may be a target of *M. hyopneumoniae* surface proteins, including cytoskeleton-associated proteins such as myosin, keratin, vimentin, and most notably for this work, actin. Actin has been found on the surface of a wide range of eukaryote cells (Chen et al., 1978; Owen et al., 1978; Jones et al., 1979; Bachvaroff et al., 1980; Rubin et al., 1982; Sanders and Craig, 1983; Rosenblatt et al., 1985b; Bach et al., 1986; Pardridge et al., 1989; Por et al., 1991; Moroianu et al., 1993; Dudani and Ganz, 1996; Smalheiser, 1996; Miles et al., 2006; Sandiford et al., 2015; Fu et al., 2017; Sudakov et al., 2017) and is a major constituent of the bronchoalveolar lavage fluid and airway surface liquid that bathes the porcine respiratory epithelium (Bartlett et al., 2013). Collectively, these observations indicate that extracellular actin may be an important receptor for *M. hyopneumoniae*.

Recent surfaceome studies indicate that 159 proteins reside on the surface of *M. hyopneumoniae* (Berry et al., 2017). Of the 143 putative actin-binding proteins identified in this study, 75 were also found on the surface of *M. hyopneumoniae* (Figure 4C)



suggesting that a number of these may have roles as adhesins. Furthermore, dominant cleavage fragments of most members of the P97 and P102 adhesin families were recovered from avidin columns coupled with biotinylated actin, including Mhp683, Mhp183, Mhp182, Mhp271, Mhp275, Mhp384, Mhp385, and Mhp493 as well as the cilium adhesin Mhp494 (P159). On a cautionary note, approximately 18% of these 143 proteins possess ATP-binding motifs. Actin monomers contain ATP and it is possible that a proportion of these actin-binding proteins are exploiting an interaction with ATP. Interestingly, many metabolic enzymes and ribosomal proteins were recovered from our affinity chromatography studies and many of these were also detected in surfaceome studies (Berry et al., 2017). These observations suggest that proteins that perform canonical functions in the cytosol have evolved to execute alternate functions on the bacterial cell surface. An excellent example of this is Elongation Factor Tu (Ef-Tu) which has recently

been shown to moonlight on the surface of *Staphylococcus aureus*, *M. hyopneumoniae*, and *Mycoplasma pneumoniae* as a multifunctional adhesin. Notably, rEf-Tu from *M. pneumoniae* bound actin with nanomolar affinity (Widjaja et al., 2017). We also identified Ef-Tu from *M. hyopneumoniae* in our affinity chromatography experiments that used actin as bait (Supplementary Table 2). Phosphoglycerate kinase (PGK) from group B streptococcus (GBS) displays four motifs ¹²⁶KKESK¹³⁰, ¹⁵⁶HRAH¹⁵⁹, ²⁰⁴KVSDK²⁰⁸, ²¹⁸KADK²²¹ that play an important role in binding actin and plasminogen (Boone and Tyrrell, 2012). *M. hyopneumoniae* binds plasminogen in a dose-dependent manner and numerous plasminogen binding proteins have been described in this species (Robinson et al., 2013; Jarocki et al., 2015; Raymond and Djordjevic, 2015). In our study, the PGK homolog from *M. hyopneumoniae* was recovered from avidin agarose columns coupled with biotinylated actin (Supplementary Table 2) and it localizes to the cell surface (Berry et al., 2017).

Three of the four actin-binding motifs in PGK from GBS are identical in the *M. hyopneumoniae* PGK homolog (data not shown). The motif KKESK has a single amino acid change to KAESK in *M. hyopneumoniae*, however the critical residue (¹³⁰K) for binding actin is conserved (Boone and Tyrrell, 2012). These data suggest that the same motifs in the *M. hyopneumoniae* PGK homolog may facilitate binding to actin and warrants further investigation. In the porcine pathogen *M. suis*, glyceraldehyde 3-phosphate dehydrogenase (GAPDH) (also known as MSP1) binds to actin on the surface of porcine red blood cells (Zhang et al., 2015). When MSG1 was expressed as a recombinant protein in *E. coli* (rMSP1) it trafficked to the cell surface and afforded *E. coli* the ability to adhere to porcine erythrocytes (Zhang et al., 2015) and affinity chromatography experiments using rMSP1 as bait identified β -actin (Zhang et al., 2015). GAPDH from group A streptococci (GAS) interacts with cytoskeletal proteins (Pancholi and Fischetti, 1992) and binds plasminogen and actin in a dose-dependent manner (Seifert et al., 2003). GAPDH is not only expressed on the surface of *M. hyopneumoniae* (Berry et al., 2017) but it was also recovered from our actin affinity chromatography experiments (Supplementary Table 2). Based on these data, surface exposed bacterial moonlighting proteins as well as more classical adhesins, such as P97, have the ability to bind to extracellular actin.

The literature is littered with descriptions of extracellular actin in a diverse array of mammalian cells (Chen et al., 1978; Owen et al., 1978; Jones et al., 1979; Bachvaroff et al., 1980; Rubin et al., 1982; Sanders and Craig, 1983; Rosenblatt et al., 1985b; Bach et al., 1986; Pardridge et al., 1989; Por et al., 1991; Moroianu et al., 1993; Dudani and Ganz, 1996; Smalheiser, 1996; Miles et al., 2006; Sandiford et al., 2015; Fu et al., 2017; Sudakov et al., 2017). Extracellular actin is likely to have evolved to be a multifunctional protein. Consistent with this view, actin is known to act as a so-called danger-associated molecular pattern during cell damage (Ahrens et al., 2012) and modulates neurotransmitter release

(Miles et al., 2006). Here we show that *M. hyopneumoniae* binds to globular β -actin and co-localizes with filamentous forms of β -actin. We have characterized a number of putative actin-binding proteins on the surface of *M. hyopneumoniae* and demonstrated that they bind actin.

AUTHOR CONTRIBUTIONS

BR conceived the confocal experiments and analyzed the 3D-SIM data. He also undertook and analyzed the protein binding and adherence assays. BR also performed the mass spectrometry experiments, analyzed the data and contributed to the drafting of the manuscript. IS prepared samples for confocal microscopy. CU cultured mycoplasma for the initial adherence assays. RM cultured mycoplasma for adherence assays, prepared samples for immunofluorescence microscopy, and aided in image analysis. MR prepared and performed scanning electron microscopy. LT and CW performed the 3D-SIM imaging experiments and assisted with interpretation of the data. MP contributed to the validation of the mass spectrometry data. SD conceived the overall study and drafted the final manuscript.

ACKNOWLEDGMENTS

This project was partly funded by the Australian Centre for Genomic Epidemiological Microbiology (Ausgem), a collaborative partnership initiated between the NSW Department of Primary Industries and the University of Technology Sydney.

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fcimb.2018.00054/full#supplementary-material>

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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