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Localization of Phosphatidylinositol 4,5-Bisphosphate to Lipid Rafts and Uroids in the Human Protozoan Parasite *Entamoeba histolytica*

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Entamoeba histolytica **is an intestinal protozoan parasite and is the causative agent of amoebiasis. During invasive infection, highly motile amoebae destroy the colonic epithelium, enter the blood circulation, and disseminate to other organs such as liver, causing liver abscess. Motility is a key factor in *E. histolytica* pathogenesis, and this process relies on a dynamic actomyosin cytoskeleton. In other systems, phosphatidylinositol 4,5-bisphosphate [$PI(4,5)P_2$] is known to regulate a wide variety of cellular functions, including signal transduction, actin remodeling, and cell motility. Little is known about the role of $PI(4,5)P_2$ in *E. histolytica* pathogenicity. In this study, we demonstrate that $PI(4,5)P_2$ is localized to cholesterol-rich microdomains, lipid rafts, and the actin-rich fractions of the *E. histolytica* membrane. Microscopy revealed that the trailing edge of polarized trophozoites, uroids, are highly enriched in lipid rafts and their constituent lipid, $PI(4,5)P_2$. Polarization and enrichment of uroids and rafts with $PI(4,5)P_2$ were enhanced upon treatment of *E. histolytica* cells with cholesterol. Exposure to cholesterol also increased intracellular calcium, which is a downstream effector of $PI(4,5)P_2$, with a concomitant increase in motility. Together, our data suggest that in *E. histolytica*, $PI(4,5)P_2$ may signal from lipid rafts and cholesterol may play a role in triggering $PI(4,5)P_2$-mediated signaling to enhance the motility of this pathogen.**

The intestinal parasite *Entamoeba histolytica* is known to cause amoebic dysentery and liver abscess. *E. histolytica* enters the human host via contaminated food or water as an environmentally stable cyst. Excystation leads to the release of trophozoites in the small intestine, which colonize the bowel lumen. From here, the parasite can enter two non-mutually exclusive routes of infection, noninvasive or invasive disease (reviewed in reference 1). In the noninvasive mode, the trophozoite encysts and exits the host, whereas in the invasive mode, the parasite adheres to and destroys the colonic epithelium and enters the circulatory system. This results in extraintestinal infection, of which amoebic liver abscess (ALA) is the most common manifestation. Motility is a key virulence function that enables this parasite to cause extraintestinal infection (2).

Motile amoebae display a polarized morphology with a pseudopod at the leading edge and a uroid (also referred to as a uropod in other eukaryotic cells) at the trailing edge. A key feature of a polarized cell is the differential distribution of proteins and lipids, including cell surface receptors, signaling molecules, and cytoskeletal elements. For example, the pseudopod of *E. histolytica* is enriched in F actin, myosin IB (3, 4), and signaling molecules like phosphatidylinositol-3,4,5-trisphosphate [$PI(3,4,5)P_3$] (5), while the uroid is enriched in myosin II and signaling molecules, including an important heterotrimeric adhesin, the galactose/*N*-acetylgalactosamine (Gal/GalNAc) lectin (6), F actin, and various actin-binding proteins (7–9). Such spatial asymmetry helps the cells to generate the necessary forces that are required for migration. For example, polymerization of actin at the leading edge facilitates pseudopod extension. Concurrently, adhesion molecules establish new focal contacts with the substrate, while activation of myosin II motors generates the contractile force required for the detachment of existing focal adhesions at the uroids. Together, these activities result in a net forward movement of the cell body. Consequently, motility in this pathogen has been attributed to its highly dynamic actomyosin cytoskeleton, which promotes the rapid morphological changes required for cell movement (reviewed in reference 10).

Phosphoinositides act as links that connect the actomyosin cytoskeleton to the plasma membrane. In other systems, phosphatidylinositol 4,5-bisphosphate [$PI(4,5)P_2$] regulates the strength of the interaction between the cytoskeleton and the plasma membrane, which is required for maintaining cell shape and integrity (11). $PI(4,5)P_2$ belongs to a group of phosphorylated phosphoinositides that act as second messenger molecules by binding to and modulating the activity of various cytoskeleton remodeling proteins. Furthermore, other lipid second messenger molecules can be generated from $PI(4,5)P_2$. For instance, the pleckstrin homology (PH) domain of phospholipase C delta 1 (PLCD1) can bind $PI(4,5)P_2$. PLCD1 hydrolyzes $PI(4,5)P_2$ to yield inositol triphosphate (IP_3) and diacylglycerol (DAG) (12, 13). While IP_3 stimulates an increase in intracellular calcium (Ca^{2+}) levels (reviewed in references 14 and 15), DAG activates protein kinase C (PKC) (reviewed in reference 16). $PI(4,5)P_2$ can be phosphorylated by phosphoinositide 3-kinases (PI3Ks) to yield yet another signaling molecule, $PI(3,4,5)P_3$ (reviewed in reference 17). All of these signaling molecules, $PI(4,5)P_2$, $PI(3,4,5)P_3$, IP_3, DAG, and Ca^{2+}, can interact with other effector proteins to trigger downstream signaling events.

In mammalian cells, $PI(4,5)P_2$ is localized to lipid rafts (18–20). Rafts are specialized membrane microdomains that are cho-

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lesterol and sphingolipid rich and are known to organize signaling molecules that participate in a variety of cellular functions, including cell polarization (21, 22), endocytosis (reviewed in reference 23), secretion (reviewed in reference 24), and adhesion (25). Previously, *E. histolytica* was shown to possess lipid rafts that regulate parasite-host interaction (26, 27). Disruption of rafts with cholesterol-binding agents like methyl-β-cyclodextrin (MβCD) inhibits adhesion of *E. histolytica* to host cells (26) and host extracellular matrix components (27). On the other hand, cholesterol loading of amoebic membranes results in enhanced adhesion to host cell monolayers, as well as cocompartmentalization of the three subunits of the Gal/GalNAc lectin in lipid rafts (28).

Whether $PI(4,5)P_2$ localizes to rafts in *E. histolytica* has not been previously studied. Therefore, in the current study, we developed a $PI(4,5)P_2$ biosensor and utilized antibody to $PI(4,5)P_2$ to study its subcellular and submembrane localization, respectively. We demonstrate that $PI(4,5)P_2$ localizes to rafts in *E. histolytica* and may signal from the raft membrane domain to regulate cell motility.

MATERIALS AND METHODS

Strains and culture conditions. *Entamoeba histolytica* trophozoites (strain HM-1:1MSS) were cultured axenically in TYI-S-33 medium (29) in 15-ml glass screw-cap tubes at 37°C.

Exposure to cholesterol or methyl-β-cyclodextrin. The source of cholesterol used was a bovine cholesterol concentrate (Creative Laboratory Products, Inc., Indianapolis, IN). In all cases, to control for extracellular sources of cholesterol, trophozoites were incubated in serum-free medium (TYI-S-33) for 30 min at 37°C prior to experimentation. After serum starvation, trophozoites were incubated in serum-free medium supplemented with 0 to 3 mg/ml cholesterol or 50 mM methyl-β-cyclodextrin (MβCD; Sigma-Aldrich, St. Louis, MO) for 15 or 60 min at 37°C. Cells treated with 0 mg/ml cholesterol represented the untreated controls for all experiments.

Lipid raft isolation, lipid extraction, and lipid dot blots. Isolation of a Triton X-100-resistant membrane and resolution of a detergent-resistant membrane (DRM) by sucrose gradient density centrifugation were carried out as previously described (26). Extraction of lipids from the sucrose gradient membrane fractions was carried out as described previously (30). Briefly, methanol–12.1 N HCl (10:1) was added to each fraction at a volumetric ratio of 1:1, and chloroform was added at a volumetric ratio of 2:1 (solvent-membrane fraction) to facilitate phase separation. The organic phase was subsequently extracted with the addition of methanol–1 N HCl (1:1) at a volumetric ratio of 1:1. The organic phase was vacuum dried, and the pellets were used for lipid dot blot analysis. Specifically, lipids were spotted onto a Hybond-C nitrocellulose membrane. The membrane was blocked in 1.5% (wt/vol) fatty acid-free bovine serum albumin (BSA) at room temperature for 1 h and probed with mouse anti-$PI(4,5)P_2$ antibody (1:1,000 dilution; Abcam, Cambridge, MA), followed by incubation with secondary antibody, peroxidase-conjugated goat anti-mouse IgG (1:2,000 dilution; Cappel, ICN Pharmaceuticals, Costa Mesa, CA). Immunoblots were visualized using an enhanced chemiluminescence Western blotting detection system (Pierce Biotechnology, Rockford, IL) according to the manufacturer's instructions. Semiquantitative densitometric analyses of immunoblots were performed using ImageJ software (version 1.42q; National Institutes of Health, Bethesda, MD). The net densitometry values were corrected to reflect the differences in membrane fraction volume. Whole-cell lipid extraction and lipid dot blot analyses were performed as previously described (31).

Generation of GST-tagged PH^{PLCD1} recombinant protein. The cDNA encoding the PH domain from PLCD1 (PH^{PLCD1}) was amplified from pEGFP-N1 (SignaGen Laboratories, Gaithersburg, MD) by PCR using the following primer pair: 5′-GGAATTCATGGACTCGGGCCGG GAC-3′ and 5′-GGTCGACTCCTTCAGGAAGTTCTGCAG-3′. The resulting PCR product was cloned into the pCR2.1-TOPO plasmid (Invitrogen, Carlsbad, CA). The DNA sequence encoding PH^{PLCD1} was excised from pCR2.1-TOPO using EcoRI and SalI endonucleases and was ligated into the polylinker region downstream of and in frame with the sequence encoding glutathione *S*-transferase (GST) in the pGEX-5X-1 expression vector (Amersham Biosciences, Piscataway, NJ). The authenticity of the DNA construct was confirmed using restriction enzyme analysis and sequencing. Recombinant GST and GST-PH^{PLCD1} proteins were obtained by expression in *Escherichia coli* BL21 (Amersham, Piscataway, NJ) according to the manufacturer's protocol. To assess the purity of the eluted proteins, SDS-PAGE and silver staining were carried out as previously described (32). To validate the GST tag, Western blotting was performed as described previously (33) using an anti-GST antibody (Chemicon, Temecula, CA) at a dilution of 1:10,000.

Protein-lipid overlay assay. A protein-lipid overlay assay was carried out to assess the $PI(4,5)P_2$ binding capability of the GST-PH^{PLCD1} fusion protein as described previously (34). Briefly, 1.6 to 100 pmol of phosphoinositides (Echelon Biosciences Inc., Salt Lake City, UT), including phosphatidylinositol 3-phosphate (PI3P), phosphatidylinositol 4-phosphate (PI4P), phosphatidylinositol 5-phosphate (PI5P), phosphatidylinositol 3,4-bisphosphate [$PI(3,4)P_2$], phosphatidylinositol 3,5-bisphosphate [$PI(3,5)P_2$], phosphatidylinositol 4,5-bisphosphate [$PI(4,5)P_2$], and phosphatidylinositol 3,4,5-triphosphate [$PI(3,4,5)P_3$], was spotted on a Hybond-C nitrocellulose membrane. The membrane was dried for 1 h at room temperature and incubated in blocking buffer containing 2% (wt/vol) fatty acid-free BSA in Tris-buffered saline (TBS)–Tween (50 mM Tris-HCl, pH 7.5, 150 mM NaCl, 0.1% [vol/vol] Tween 20) at room temperature for 1 h. The membrane was incubated overnight at 4°C in blocking buffer containing 0.2 μg/ml of either GST or GST-PH^{PLCD1}, washed in TBS-Tween, and incubated in polyclonal anti-GST antibody (1:10,000 dilution) in blocking buffer for 1 h. The membrane was washed, incubated in horseradish peroxidase (HRP)–anti-rabbit IgG (1:5,000 dilution) in TBS-Tween for 1 h, and then washed and developed as described previously (33).

We also tested the ability of GST or GST-PH^{PLCD1} to bind *E. histolytica* cell extracts. Whole-cell protein extracts of trophozoites (5×10^5 and 1×10^5 cells) were resolved by SDS-PAGE and electroblotted onto a nitrocellulose membrane. The membrane was washed twice with distilled water and dried for 1 h at room temperature. Serial dilutions of $PI(4,5)P_2$ were spotted on one edge of the same membrane, which served as a positive control for binding of the probe. The membrane was dried and incubated with blocking buffer containing 2% (wt/vol) fatty acid-free BSA in TBS-Tween (50 mM Tris-HCl, pH 7.5, 150 mM NaCl, 0.1% [vol/vol] Tween 20), followed by incubation with 0.2 μg/ml of either GST or GST-PH^{PLCD1} probes in the blocking buffer, as described above. The membrane was washed in TBS-Tween and incubated in polyclonal anti-GST antibody (1:10,000 dilution) in blocking buffer, followed by incubation with HRP–anti-rabbit IgG (1:5,000 dilution) in TBS-Tween, and then washed and developed as described previously (33). As a positive control for protein transfer, the membrane was incubated in anti-actin monoclonal antibody (1:2,500 dilution; Abcam) for 1 h. The membrane was washed, incubated in HRP–anti-mouse IgG (1:2,000 dilution) in TBS-Tween for 1 h, and then washed and developed as described previously (33).

Biosensor staining and fluorescence microscopy. For fluorescence microscopy, *E. histolytica* trophozoites were plated on a chambered cover glass slide system (Lab-Tek, Christchurch, New Zealand) and fixed with 4% (vol/vol) paraformaldehyde overnight at 37°C. Cells were washed in phosphate-buffered saline (PBS), incubated in 50 mM NH_4Cl in PBS for 15 min, and then permeabilized with 0.05% (wt/vol in PBS) saponin for 10 min as previously described (35). All subsequent buffers contained 0.05% (wt/vol in PBS) saponin. This regimen preserves the cellular morphology and permeabilizes cells without removing phosphoinositides from the plasma membrane (5, 35). Cells were blocked with 10% (vol/vol) fetal bovine serum (FBS)–PBS and stained with approximately 10 μg/ml

of either affinity-purified GST or GST-PH^{PLCD1}, followed by staining with Alexa Fluor 488-conjugated rabbit anti-GST antibody (1:2,000; Invitrogen) or Texas Red-conjugated goat anti-GST antibody (1:1,000; Rockland Immunochemicals, Gilbertsville, PA), as described previously (35). Stained cells were viewed using a Nikon Eclipse TI-E spectral confocal microscope (Nikon Instruments Inc., Lewisville, TX) or an LSM510 confocal microscope (Carl Zeiss MicroImaging, Thornwood, NY). Images were analyzed using Nikon NIS Elements or LSM Image Browser software, respectively. To assess the effect of cholesterol on localization of $PI(4,5)P_2$, cells were exposed to either 0 or 3 mg/ml cholesterol for 1 h and then fixed and stained as described above.

For lipid raft staining, *E. histolytica* trophozoites were fixed with 4% (vol/vol) paraformaldehyde overnight at 37°C. Dialkyindocarbocyanine ($DiIC_{16}$; Invitrogen) staining of lipid rafts was carried out as described previously (27).

To determine if $PI(4,5)P_2$ and lipid rafts localized to uroids, capping was induced by incubation of amoebae with Alexa Fluor 488-conjugated concanavalin A (ConA; Invitrogen) for 1 h on ice, followed by incubation on cover glass chambered slides for 15 min at 37°C as described previously (6). $PI(4,5)P_2$ or lipid raft staining was carried out as described above.

Measurement of motility. To test the effect of cholesterol on parasite motility, a motility assay was employed according to the methods of Zaki et al. (36). Briefly, cells were serum starved for 30 min at 37°C and exposed to a range of cholesterol concentrations (0 to 3 mg/ml) in serum-free medium for 30 min at 37°C. Eight milliliters of complete TYI-S-33 medium supplemented with 0.75% (wt/vol) agarose was poured into 60-mm petri dishes and allowed to solidify at room temperature. A trough (2 by 30 mm) was cut into the solidified medium to serve as the motility chamber. Following exposure to cholesterol, trophozoites (5×10^5) were placed in the trough of the motility chamber. A coverslip (22 by 40 mm) was placed over the trough and the plate was incubated at 37°C for 3 h in 5% CO_2. Images were captured using confocal microscopy. The migration distance was measured using Zeiss LSM510 image analysis software.

Measurement of intracellular Ca^{2+} levels. To measure the effect of cholesterol treatment on intracellular Ca^{2+} levels, a Ca^{2+} assay was employed as described previously (31). Briefly, cells were incubated in serum-free medium supplemented with the fluorescent Ca^{2+} indicator Fluo-4/AM (Invitrogen) or dimethyl sulfoxide (DMSO; diluent control) for 30 min at 37°C. Then, cells were incubated with a range of concentrations of cholesterol (0 to 3 mg/ml) for either 15 min or 1 h at 37°C. Subsequently, 1×10^5 trophozoites were pipetted into the wells of an uncoated 96-well plate, and after 5 min the plates were transferred to a fluorescence microplate reader for 30 min. To account for the background fluorescence, the fluorescence values for cells incubated in DMSO were subtracted from the fluorescence values for cells incubated in Fluo-4/AM. The fluorescence value of the untreated control cells at the 35-min time point was arbitrarily set equal to 100%, and the values at other time points were reported as a percentage of this value.

Statistical analyses. All values are given as means ± standard deviations (SDs) of at least 3 trials. To compare means, statistical analyses were performed using GraphPad Instat v.3 software with an unpaired *t* test, Welch corrected (two-tail *P* value), or with one-way analysis of variance (ANOVA) and a Tukey-Kramer multiple-comparison test. In all cases, *P* values of less than 0.001 were considered highly statistically significant, while *P* values of less than 0.01 or 0.05 were considered statistically significant.

RESULTS

$PI(4,5)P_2$ compartmentalizes in lipid rafts and actin-rich pellet fractions of the DRM in *E. histolytica*. In mammalian cells, $PI(4,5)P_2$ has been shown to localize to lipid rafts (18, 19). Therefore, we determined if *E. histolytica* exhibits similar compartmentalization of $PI(4,5)P_2$. The detergent-resistant membrane (DRM), which consists of lipid rafts and the actin-rich membrane, was isolated as described previously (26). Sucrose gradient density centrifugation was used to resolve these two membrane types. Total phosphoinositides were extracted from each of the gradient fractions and subjected to a lipid dot blot analysis using antibodies to $PI(4,5)P_2$ (Fig. 1A). We previously demonstrated that membrane fractions 9 to 14 possess high levels of cholesterol and contain lipid rafts (31). Less buoyant fractions (fractions 17 to 19) and the pellet fraction (fraction 20) are actin rich (26, 28, 31). Similar to the enrichment seen in mammalian cells, we observed an enrichment of $PI(4,5)P_2$ in lipid rafts (fractions 9 to 14) (Fig. 1A and B). However, the actin-rich pellet fraction also possessed high levels of this phospholipid (Fig. 1A and B). Thus, there may be two pools of $PI(4,5)P_2$ in *E. histolytica*, one pool in lipid rafts and another pool interacting directly with the actin cytoskeleton. However, we cannot rule out the possibility that $PI(4,5)P_2$ in rafts also interacts with the actin cytoskeleton in live cells.

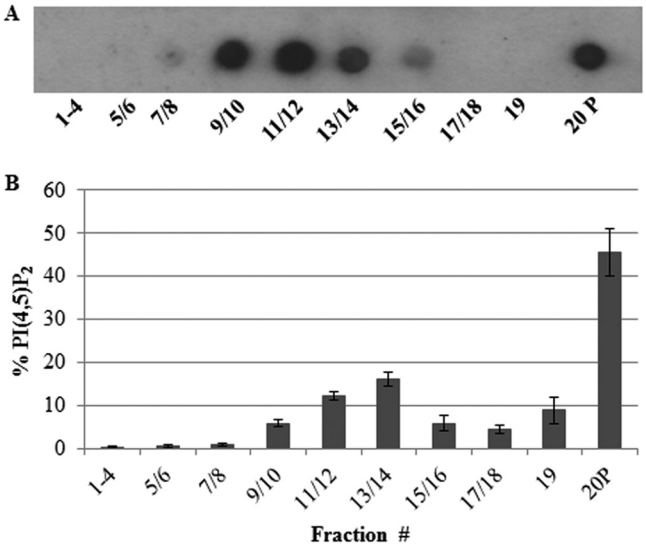

FIG 1 $PI(4,5)P_2$ compartmentalizes in lipid rafts and actin-rich pellet fractions. The DRM was isolated from *Entamoeba histolytica* trophozoites and subjected to sucrose gradient density fractionation. $PI(4,5)P_2$ was extracted from these fractions, and lipid dot blots were performed using antibodies specific to $PI(4,5)P_2$. (A) A representative blot is shown. Cells displayed $PI(4,5)P_2$ enrichment in lipid raft fractions (fractions 9 to 14) and in an actin-rich pellet fraction (fraction 20 P). (B) Semiquantitative scanning densitometry was used to analyze the distribution of $PI(4,5)P_2$ in membrane fractions, correcting for the difference in the starting volume of each membrane fraction. The total level of $PI(4,5)P_2$ in the DRM was calculated, and the percentage of $PI(4,5)P_2$ in each fraction (± SD; $n = 3$) is presented.

Exposure to cholesterol increases levels of $PI(4,5)P_2$ in the DRM. Previous studies in mammalian cells have revealed that the level of membrane cholesterol influences the localization of $PI(4,5)P_2$ in the DRM (18). Therefore, we tested the effect of exposure to cholesterol or to the cholesterol-sequestering agent MβCD on the localization and overall levels of $PI(4,5)P_2$ in *E. histolytica* trophozoites. On average, cells treated with 0.05 mg/ml cholesterol displayed a slight increase in $PI(4,5)P_2$ levels in the lipid raft fractions compared to control untreated cells; however, there was no difference in the level of $PI(4,5)P_2$ in the total DRM (Fig. 2A to C). The most dramatic changes in the level of $PI(4,5)P_2$ in the total DRM were observed when the cells were exposed to 0.75 and 3 mg/ml cholesterol, with an approximately 60% increase (Fig. 2A). In contrast, MβCD caused a 56% decrease in the

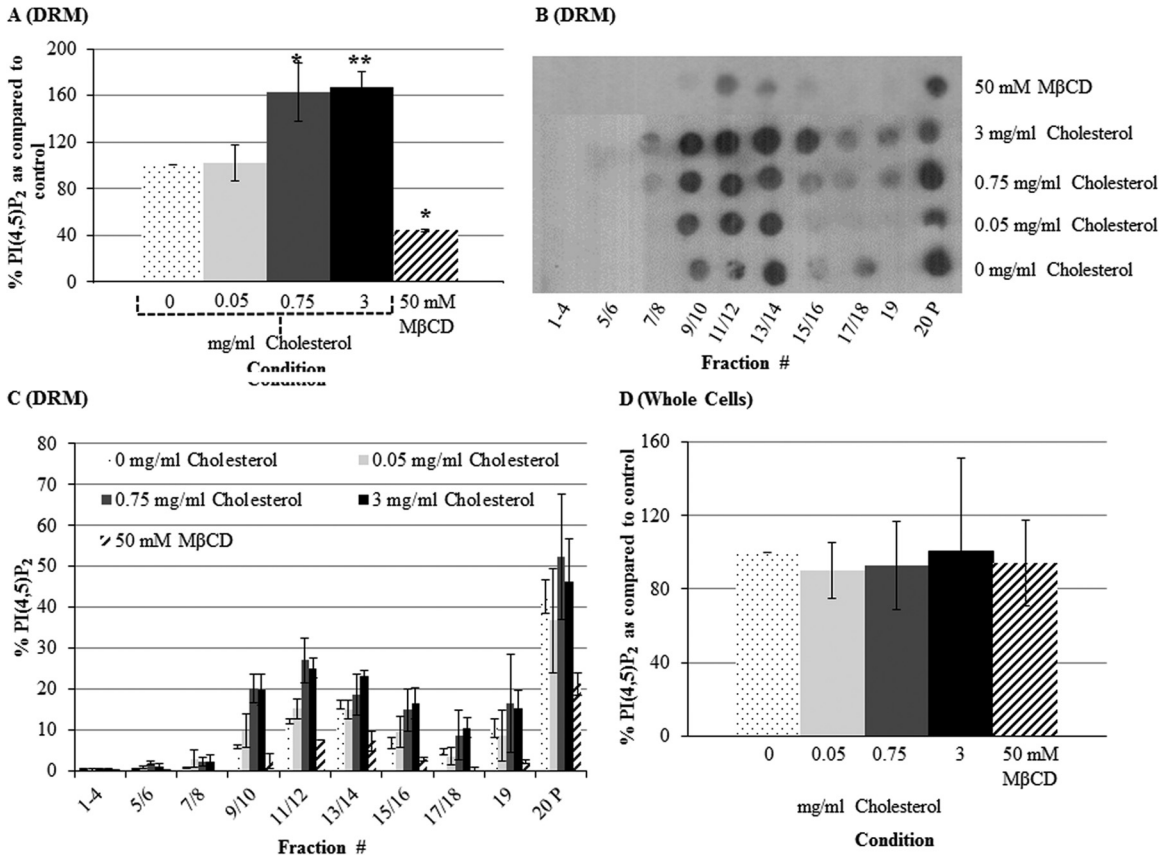

FIG 2 Exposure to cholesterol increases the levels of $PI(4,5)P_2$ in the DRM and specifically in lipid raft fractions. Trophozoites were serum starved and exposed to cholesterol (0 to 3 mg/ml) or the cholesterol-depleting agent MβCD. Cells treated with no cholesterol (0 mg/ml) served as the untreated control. (A to C) The DRM was isolated and subjected to sucrose gradient density fractionation. $PI(4,5)P_2$ was extracted from these fractions, and lipid dot blots were performed using antibodies specific to $PI(4,5)P_2$. Levels of $PI(4,5)P_2$ were analyzed by semiquantitative scanning densitometry. In all cases, values were corrected for differences in starting volumes. (A) The total level of $PI(4,5)P_2$ in the DRM was calculated and reported as a percentage of that for the untreated control. Cells treated with cholesterol (0.75 or 3 mg/ml) and MβCD displayed an approximately 60% increase and 56% decrease in $PI(4,5)P_2$ in the total DRM, respectively, compared to the levels for the control ($n \geq 3$) (*, $P < 0.5$; **, $P < 0.01$). (B) A representative lipid dot blot of the sucrose gradient density fraction is shown. Cells treated with cholesterol (0.75 and 3 mg/ml) displayed enhanced enrichment of $PI(4,5)P_2$ in lipid rafts (fractions 9 to 14) compared to the control, while MβCD-treated cells showed a decrease in $PI(4,5)P_2$ from all fractions of the DRM. (C) The data represent the percentage of $PI(4,5)P_2$ in each fraction compared to the level of $PI(4,5)P_2$ in the total DRM of the untreated control cells (± SD; $n \geq 3$). (D) Phosphoinositides were extracted from whole-cell lysates, and $PI(4,5)P_2$ levels were measured using dot blots with antibodies specific to $PI(4,5)P_2$. The level of $PI(4,5)P_2$ for each treatment was quantified by scanning densitometry and reported as a percentage of the level for the untreated control. The total cellular $PI(4,5)P_2$ level remained unchanged upon treatment with cholesterol or MβCD. Data are the means ± SDs for ≥3 independent experiments.

level of $PI(4,5)P_2$ from that in the total DRM (Fig. 2A). Furthermore, treatment with 0.75 and 3 mg/ml cholesterol increased the level of $PI(4,5)P_2$ specifically in lipid rafts (Fig. 2B and C). To determine if these changes were due to overall increases or decreases in $PI(4,5)P_2$, we also quantified total $PI(4,5)P_2$ in whole cells. We did not find a significant difference in the total cellular level of $PI(4,5)P_2$ after cholesterol or MβCD exposure (Fig. 2D). Thus, $PI(4,5)P_2$ may be moving between membrane compartments in a cholesterol-dependent manner. However, we cannot rule out the possibility that there is a rapid turnover of $PI(4,5)P_2$, where continuous synthesis and hydrolysis maintain a constant total level.

A GST-tagged PH probe reveals localization of $PI(4,5)P_2$ to plasma membrane and uroids. We have previously determined the subcellular localization of PI3P (35) and $PI(3,4,5)P_3$ (5) using a GST-tagged FYVE finger domain and a GST-tagged pleckstrin homology (PH) domain of Bruton's tyrosine kinase (PH^{BTK}), respectively. We used a similar approach to determine the subcellular localization of $PI(4,5)P_2$ using a recombinant GST-tagged version of the PH domain from PLCD1 (PH^{PLCD1}). This protein domain has been shown to specifically bind $PI(4,5)P_2$, and the GST-tagged chimera has been used to study the localization of $PI(4,5)P_2$ in mammalian cells (37).

Recombinant GST (control) and GST-PH^{PLCD1} were expressed in bacteria and affinity purified. SDS-PAGE demonstrated that the proteins were 28 kDa and 48 kDa in size, which are consistent with the predicted sizes of GST and GST-PH^{PLCD1}, respectively (Fig. 3A). The authenticity of the GST tag was confirmed by Western blotting with antibodies specific for GST (Fig. 3B). To test the binding specificity of GST-PH^{PLCD1}, we employed a protein-lipid overlay assay as described previously (34, 37). The probe bound to $PI(4,5)P_2$ in a dose-dependent manner and did not display any significant capacity for binding to other related phosphoinositides (Fig. 3C). GST alone (control) did not bind any of

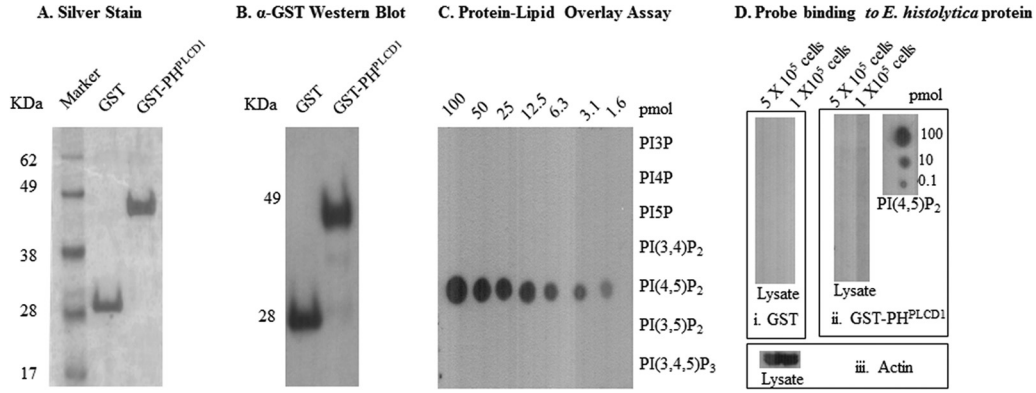

FIG 3 The purified GST-PH^{PLCD1} biosensor binds with high affinity to PI(4,5)P_2. (A) SDS-PAGE and silver staining of GST and GST-PH^{PLCD1} fusion proteins after affinity purification. The molecular mass markers are shown. (B) Western blot analysis of purified GST and GST-PH^{PLCD1} using anti-GST antibody. (C) Serial dilutions (100 to 1.6 pmol) of the indicated phosphoinositides were spotted onto a nitrocellulose membrane and incubated with 0.2 μg/ml of GST-PH^{PLCD1}. Binding of the protein was detected with anti-GST antibodies. The probe exhibited binding specificity to PI(4,5)P_2 alone. (D) Whole-trophozoite extracts (5×10^5 and 1×10^5 cells) were resolved by SDS-PAGE and electroblotted onto a Hybond-C nitrocellulose membrane. Individual lanes of lysate were decorated with GST (i), GST-PH^{PLCD1} (ii), or anti-actin antibody (positive control for protein transfer) (iii). (ii) As an additional control for binding of the GST-PH^{PLCD1} probe, the membrane was washed after the protein transfer and serial dilutions of PI(4,5)P_2 were spotted on the same membrane and incubated with GST-PH^{PLCD1}. Binding of the probes was detected with anti-GST antibodies. Neither GST nor GST-PH^{PLCD1} probe bound to the whole-cell protein extract, further supporting the specificity of the probe.

the phosphoinositides (data not shown). We also determined if the probes bound nonspecifically to *E. histolytica* proteins. Whole-cell protein extract was resolved by SDS-PAGE, electroblotted onto a nitrocellulose membrane, and decorated with GST or GST-PH^{PLCD1}. Neither GST nor GST-PH^{PLCD1} displayed binding to *E. histolytica* proteins (Fig. 3D). Together, these data suggest that the GST-PH^{PLCD1} probe is highly specific for PI(4,5)P_2 and does not cross-react with other cellular lipids or proteins. Therefore, it could be utilized for subcellular localization of PI(4,5)P_2.

E. histolytica trophozoites were fixed, permeabilized, and stained with GST or GST-PH^{PLCD1}, followed by staining with a fluorescence-labeled anti-GST antibody. Confocal microscopy revealed that cells stained with control GST exhibited minimal fluorescence (Fig. 4A and D). In contrast, the GST-PH^{PLCD1} probe uniformly decorated the plasma membrane, giving most cells a ring-like appearance (Fig. 4B and E). This was not surprising, as PI(4,5)P_2 is known to account for 1% of plasma membrane phospholipids and >99% of all doubly phosphorylated phospholipids and is maintained at a high steady-state level in mammalian cells (reviewed in references 17, 20, and 38). Interestingly, in some cells, PI(4,5)P_2 was also enriched on one side of the cell, opposite apparent forming pseudopods (Fig. 4C and F).

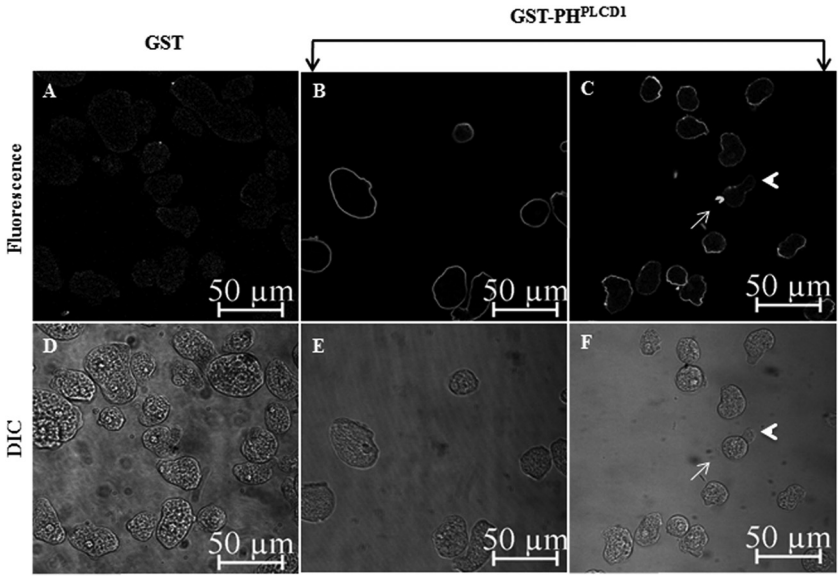

FIG 4 The PI(4,5)P_2 biosensor GST-PH^{PLCD1} decorates the plasma membrane of *E. histolytica*. Trophozoites were stained with GST or GST-PH^{PLCD1} and fluorescent anti-GST antibody, and immunofluorescence confocal microscopy was performed. (A) Cells stained with the control, GST, exhibited minimal staining. (B) GST-PH^{PLCD1} uniformly decorated the plasma membrane of the trophozoites, giving the cells a ring-like appearance. (C) In some instances, there was an enrichment of GST-PH^{PLCD1}staining on one edge of the cell (arrow), opposite an apparent pseudopod (arrowhead). (D to F) The differential interference contrast (DIC) images corresponding to the images in panels A to C, respectively, are shown.

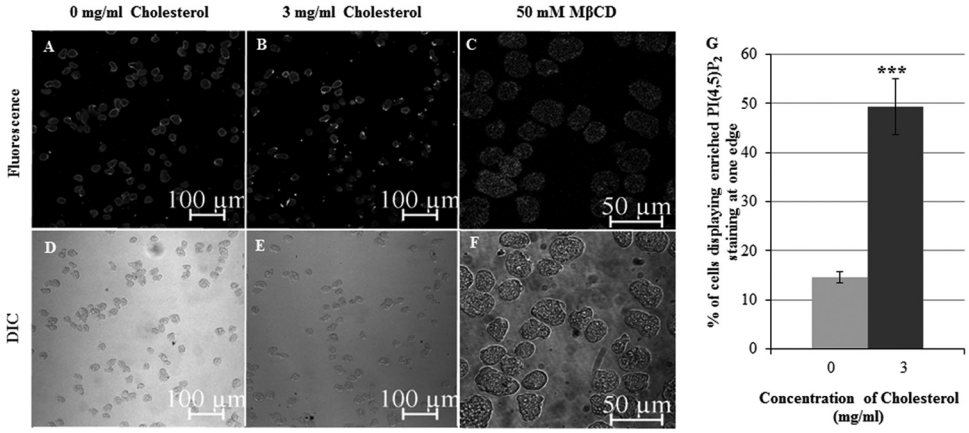

FIG 5 Treatment with cholesterol enhances the number of cells displaying $PI(4,5)P_2$ enrichment at the edge opposite the forming pseudopod. Trophozoites were serum starved and exposed to either 0 mg/ml cholesterol (control), 3 mg/ml cholesterol, or the lipid raft-disrupting agent MβCD. Cells were stained with GST-PH^{PLCD1} and fluorescent anti-GST antibody, and immunofluorescence confocal microscopy was performed. (A and B) Cholesterol-treated cells displayed an increased $PI(4,5)P_2$ localization on one edge of the cell compared to untreated control cells. (C) In MβCD-treated cells, $PI(4,5)P_2$ staining was no longer confined to the plasma membrane. (D to F) The DIC images corresponding to the images in panels A to C, respectively, are shown. (G) The percentage of apparently polarized cells in control cells or after cholesterol treatment was quantified by visually scoring cells at ×20 magnification. There was a 3.4-fold increase in the number of cells displaying $PI(4,5)P_2$ localization on one edge after cholesterol treatment compared to the number for untreated control cells. Data are the means ± SDs of 3 independent experiments (***, $P < 0.001$).

Studies in other systems (18) and our biochemical analysis (Fig. 2A to C) indicated that cholesterol levels impact the localization of $PI(4,5)P_2$ in the DRM. Therefore, we wanted to determine the subcellular localization of $PI(4,5)P_2$ in whole cells after exposure to cholesterol or after cholesterol depletion. Interestingly, there was an increase in the number of cells displaying $PI(4,5)P_2$ localization at one edge upon cholesterol treatment (Fig. 5B and E) compared to the number for the untreated control (Fig. 5A and D). Image analysis revealed that there was a 3.4-fold increase in the number of cells displaying this morphology after cholesterol treatment compared to the number of control cells (Fig. 5G). In contrast, cholesterol depletion with MβCD disrupted the localization of $PI(4,5)P_2$ at the plasma membrane, in that it was observed in the interior of the cell (Fig. 5C and F). This suggested a requirement of cholesterol richness for proper localization of $PI(4,5)P_2$ to the plasma membrane.

Cholesterol treatment enhances motility. In other systems, $PI(4,5)P_2$ plays an essential role in cell motility. Since $PI(4,5)P_2$ became asymmetrically distributed after exposure to cholesterol (Fig. 5B and E) and since asymmetry is a hallmark of motile cells, we hypothesized that exposure to cholesterol was enhancing motility. To test this, we utilized a previously published under-agar motility assay (36) to determine the effect of cholesterol exposure on cell movement. Upon exposure to cholesterol, the ability of the trophozoites to move out of the wells increased significantly in a dose-dependent fashion (Fig. 6A). After exposure to cholesterol, not only did cells traverse longer distances but also there was an apparent increase in the number of motile cells (Fig. 6B and C). This suggests a possible role for cholesterol in mediating motility in *E. histolytica*.

$PI(4,5)P_2$ localizes to uroids in ConA-capped *E. histolytica* cells. In apparently polarized cells, $PI(4,5)P_2$ appeared to be enriched on one edge opposite that of the forming pseudopods. This could represent localization of $PI(4,5)P_2$ to the trailing edge (uroid). Therefore, we employed a polyvalent ligand, ConA, which is known to interact with and cap cell surface receptors at the trailing edge of the trophozoites (39). Thus, labeled ConA serves as a uroid marker. $PI(4,5)P_2$ colocalized with fluorescent ConA in uroids (Fig. 7A to D) and with patches of ConA in trophozoites in which capping was initiated but was not complete (data not shown).

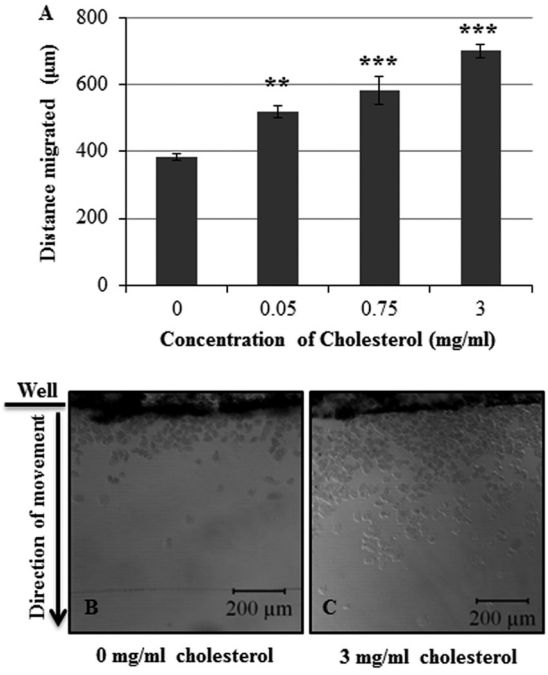

FIG 6 Exposure to cholesterol enhances motility in *E. histolytica* cells. *E. histolytica* trophozoites were serum starved and exposed to 0 to 3 mg/ml cholesterol. A motility assay was performed, and images were captured using confocal microscopy. The distance migrated by trophozoites was measured. (A) Trophozoites exhibited a dose-dependent and statistically significant increase in motility after exposure to cholesterol (**, $P < 0.01$; ***, $P < 0.001$). The data represent the means ± SDs from ≥3 independent experiments. (B and C) Cholesterol treatment also increased the number of motile cells compared to the number of untreated control cells entering the agar. Representative ×10 magnification fields of control cells and cells treated with 3 mg/ml cholesterol are shown.

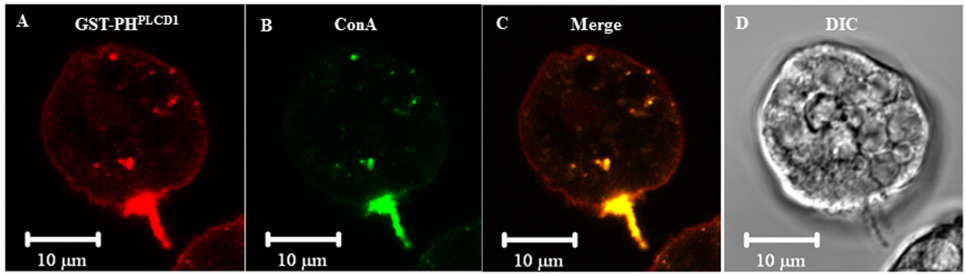

FIG 7 Fluorescence confocal microscopy of ConA-capped *E. histolytica* reveals localization of PI(4,5)P_2 to uroids. (A to D) Trophozoites were incubated with fluorescein isothiocyanate-ConA (green) for 1 h on ice, followed by incubation at 37°C for 15 min to initiate uroid formation. Cells were fixed and stained with GST-PH^{PLCD1} and anti-GST antibody (red).

Lipid rafts localize to uroids. Since our biochemical analysis demonstrated that lipid rafts were enriched in PI(4,5)P_2 and since microscopy revealed that PI(4,5)P_2 localized to uroids, we wanted to determine if, like in mammalian cells (40), the uroids of *E. histolytica* were enriched in lipid rafts. Therefore, we employed a lipid raft stain, DiIC$_{16}$, on fixed *E. histolytica* trophozoites as previously described (27). Cells with no apparent pseudopod formation (nonpolarized) displayed a uniform plasma membrane raft staining (Fig. 8A and B), whereas apparently polarized cells also showed an enrichment of raft staining at one edge opposite the forming pseudopod (Fig. 8C and D). Colocalization of DiIC$_{16}$ with ConA revealed that the uroids were indeed enriched in lipid rafts (Fig. 8E to H). Similar to the results seen for PI(4,5)P_2 staining, trophozoites with incomplete uroid formation exhibited DiIC$_{16}$ staining that colocalized with lateral patches of ConA (data not shown).

Cholesterol treatment enhances intracellular Ca^{2+} levels in *E. histolytica*. In other systems, PLCD1 hydrolyzes PI(4,5)P_2 into IP_3 and DAG, which facilitates Ca^{2+} signaling (13, 41). Since PLCD1 also localizes to rafts (42), it is reasonable to hypothesize that the accumulation of PI(4,5)P_2 in rafts after cholesterol treatment might lead to an increase in intracellular Ca^{2+}. Therefore, we measured intracellular Ca^{2+} levels after cholesterol treatment using a previously published protocol (31). Several studies have demonstrated that bona fide ligands of *E. histolytica*, such as collagen and fibronectin themselves, induce signaling leading to in-

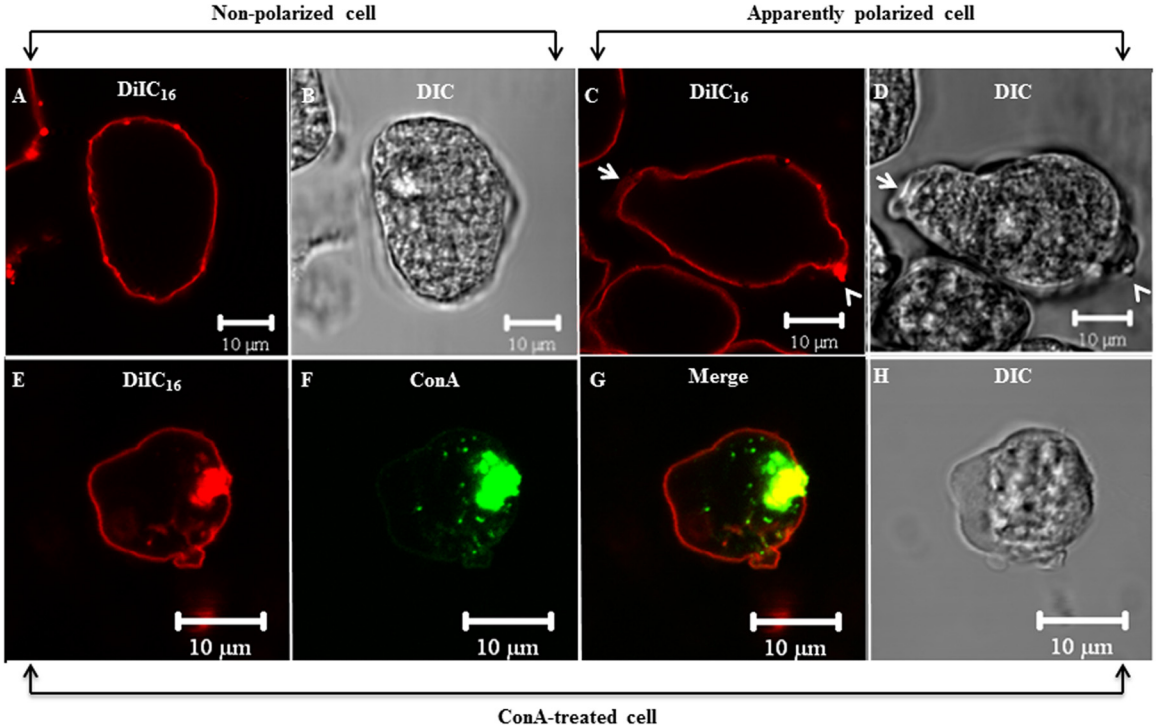

FIG 8 Fluorescence confocal microscopy of ConA-capped *E. histolytica* reveals localization of lipid rafts to uroids. The lipid raft localizes to uroids in *E. histolytica*. (A to D) Trophozoites were fixed and stained with the lipid raft marker DiIC$_{16}$. Nonpolarized cells displayed uniform plasma membrane raft staining, while apparent polarized cells exhibited an enriched raft staining on one edge (arrowheads) opposite the pseudopod (arrows). (E to H) *E. histolytica* trophozoites were incubated with fluorescein isothiocyanate-ConA (green) for 1 h on ice, followed by incubation at 37°C for 15 min to initiate uroid formation. Cells were fixed and stained with the raft marker DiIC$_{16}$ (red). There was a colocalization of ConA and DiIC$_{16}$ staining at the uroid of *E. histolytica*, further supporting the finding that lipid rafts are enriched at uroids in polarized cells.

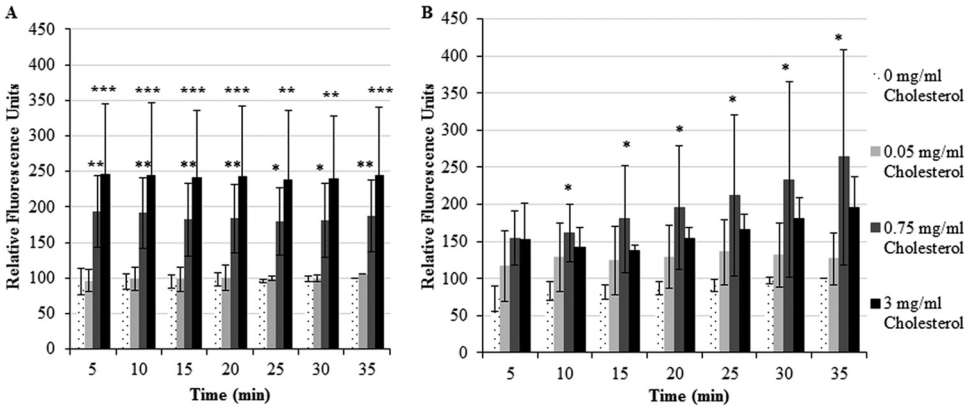

FIG 9 Exposure to cholesterol increases intracellular Ca^{2+} levels in *E. histolytica*. Trophozoites were serum starved and exposed to cholesterol at concentrations of 0 (control) to 3 mg/ml for either 60 or 15 min. Intracellular Ca^{2+} levels were measured after treatment over a 35-min time interval. Intracellular Ca^{2+} levels at the 35-min time point for the untreated control cells were arbitrarily set equal to 100%, and the values at the other time points were reported as a percentage of this final value. The data are the means ± SDs for ≥3 independent experiments. (A) Sixty minutes of pretreatment with 0.75 or 3 mg/ml cholesterol resulted in a significant increase in the intracellular Ca^{2+} level compared to that for untreated controls (*, $P < 0.5$; **, $P < 0.01$; ***, $P < 0.001$). (B) Similarly, an increase in intracellular Ca^{2+} levels was observed with a 15-min cholesterol pretreatment (*, $P < 0.5$).

creased intracellular Ca^{2+} (31, 43). Therefore, to test the effect of cholesterol alone on intracellular Ca^{2+} levels, we utilized uncoated 96-well plates for our studies. After 1 h of pretreatment with cholesterol, we observed a significant dose-dependent increase in Ca^{2+} (Fig. 9A). Since signaling events take place immediately following stimulation, we also tested a shorter time of exposure to cholesterol. After 15 min of pretreatment with 0.75 and 3 mg/ml cholesterol, we observed similar increases in Ca^{2+}, of which some were statistically significant (Fig. 9B). Together, these data indicate the possibility of a cholesterol-dependent, $PI(4,5)P_2$-mediated rise in intracellular Ca^{2+} levels.

DISCUSSION

This study represents the first analysis of the submembrane localization of $PI(4,5)P_2$ in *E. histolytica*. Biochemical analyses showed that $PI(4,5)P_2$ is compartmentalized in lipid raft and actin-rich pellet fractions of the DRM. Microscopy demonstrated that $PI(4,5)P_2$ uniformly decorated the plasma membrane in a majority of the cells but also colocalized with lipid rafts at the uroids in polarized cells. The connection between $PI(4,5)P_2$ and rafts in *E. histolytica* was supported by our observations that cholesterol treatment increased the level of $PI(4,5)P_2$ in the DRM, lipid rafts, and raft-rich uroids and that MβCD depleted the DRM and plasma membrane of $PI(4,5)P_2$. Finally, treatment with cholesterol resulted in an increase in intracellular Ca^{2+} [a downstream effector of $PI(4,5)P_2$] and a concomitant increase in motility. Therefore, in *E. histolytica*, $PI(4,5)P_2$ may signal from rafts and cholesterol may play a role in triggering $PI(4,5)P_2$-dependent signaling.

Various studies have supported the existence of different cellular pools of $PI(4,5)P_2$ to explain the complex roles of $PI(4,5)P_2$ in cells (reviewed in reference 38). In *E. histolytica*, we observed compartmentalization of $PI(4,5)P_2$ in lipid rafts as well as in the actin-rich pellet fraction at steady state. We also showed that exposure to cholesterol resulted in enrichment of $PI(4,5)P_2$ in the DRM and, specifically, in lipid rafts without an increase in the total cellular level of $PI(4,5)P_2$. One explanation for this is that $PI(4,5)P_2$ translocates between membrane subcompartments in a cholesterol-dependent fashion. The parasite likely encounters different extracellular cholesterol levels depending on the location in the human host (e.g., liver versus intestine). Therefore, simple translocation of $PI(4,5)P_2$ between membrane domains would represent a rapid way to regulate lipid signaling throughout infection. Ca^{2+} is a downstream effector of $PI(4,5)P_2$ and accumulates upon hydrolysis of $PI(4,5)P_2$ to IP_3 and DAG. Since we also observed an increase in the intracellular Ca^{2+} level upon cholesterol treatment, it is also possible that there is rapid hydrolysis and synthesis of $PI(4,5)P_2$ in rafts, which would maintain the apparent constant total cellular level of this lipid. In other systems, in addition to PLCD1 (42), enzymes that synthesize $PI(4,5)P_2$, such as phosphatidylinositol-4-phosphate-5-kinase (PIP5K), are also recruited to rafts upon stimulation (44, 45). Thus, it is conceivable that rapid turnover of $PI(4,5)P_2$ could occur in rafts.

We cannot rule out the possibility that it is actually the rise in intracellular Ca^{2+} that induces segregation of $PI(4,5)P_2$ into rafts. Using an *in vitro* lipid monolayer system, Levental et al. (46) showed that the lateral organization of $PI(4,5)P_2$ is sensitive to small changes in Ca^{2+} concentration. However, it is not known if Ca^{2+} can influence the submembrane distribution of $PI(4,5)P_2$ *in vivo*. Regardless of whether there is a Ca^{2+}-dependent step, the outcome of exposure of trophozoites to cholesterol is enrichment of $PI(4,5)P_2$ in lipid rafts. This provides significant insight into signaling in this parasite.

In the present study, we demonstrated that $PI(4,5)P_2$ uniformly decorates the plasma membrane of this parasite. We also showed that in polarized cells, both lipid rafts and the constituent signaling lipid, $PI(4,5)P_2$, localize to the uroids. In mammalian cells, there is increasing evidence that lipid rafts play a vital role in delivering signaling molecules to uroids during cell migration. For example, Millán et al. (40) showed that the uroids of polarized T lymphocytes are enriched in cholesterol and disruption of rafts leads to loss of uroids and subsequent inhibition of cell migration. Furthermore, segregation of membrane proteins to uroids and pseudopods is inhibited by disruption of lipid rafts in T lymphocytes (40, 47) and neutrophils (21). Another study of polarized neutrophils revealed that $DiIC_{16}$-stained raft fractions progress from a uniform distribution to discrete patches that coalesce at

uroids to form caps (48). Furthermore, cap formation in these cells relies on the interaction of transmembrane proteins of the DRM with both the lipids of the DRM and the proteins of the actomyosin cytoskeleton (48). Since we observed the localization of lipid rafts and PI(4,5)P_2 in uroids and in ConA-positive, laterally localized membrane patches, it is possible that a similar lipid raft-mediated delivery of PI(4,5)P_2 to uroids occurs in *E. histolytica*. Another mechanism by which PI(4,5)P_2 could become enriched in uroids is by local synthesis. A number of reports in other systems have shown that key PI(4,5)P_2-synthesizing enzymes also localize to uroids (49–52). However, whether these enzymes localize to uroids of *E. histolytica* remains to be determined.

PI(4,5)P_2 can function as a second messenger and bind various actin-binding proteins. A proteomic analysis of *E. histolytica* uroids identified the presence of actin-binding proteins that have been reported to interact with PI(4,5)P_2 in other eukaryotic cells, including the spectrin-family proteins (e.g., filamin 2, an actinin-like protein) and a novel protein, filopodin (53). In other systems, PI(4,5)P_2 has been shown to regulate actin cytoskeleton by binding and modulating filamin (54) and α-actinin (55). Filopodin is an ezrin-radixin-moesin (ERM) domain-containing protein. Members of the ERM family bind PI(4,5)P_2 and are known to regulate uroid formation in T cells (56). Thus, PI(4,5)P_2 may serve as a bridge at the uroid, connecting the cytoskeleton to the plasma membrane through various actin-binding proteins.

PI(4,5)P_2 can also generate a variety of secondary messenger molecules. For instance, at the leading edge of *Dictyostelium discoideum* cells, PI(4,5)P_2 is converted to PI(3,4,5)P_3 by PI3K; PI(3,4,5)P_3 binds to and recruits several proteins to promote pseudopod formation (reviewed in reference 52). Likewise, PI(3,4,5)P_3 localizes to pseudopods of motile *E. histolytica* (5). At the trailing edge of mammalian cells, hydrolysis of PI(4,5)P_2 to DAG, which activates PKC, is essential for T cell chemotaxis (57). IP_3, another product of PI(4,5)P_2 hydrolysis, stimulates an increase in intracellular Ca^{2+} levels, and increases in Ca^{2+} correlate with enhanced motility in neutrophils (58, 59). Ca^{2+} is a potent stimulator of motility through activation of myosin light-chain kinase (MLCK), leading to an increase in myosin II-based contractility at the uroid (60). In the current study, an increase in intracellular Ca^{2+} was accompanied by a robust augmentation of motility after exposure to cholesterol. Thus, a similarly potent PI(4,5)P_2 to Ca^{2+} to MLCK system may exist in this pathogen. In support of this, myosin II is found to localize to the uroids of *E. histolytica* (2).

On average, we observed a modest increase in PI(4,5)P_2 levels in rafts after treatment with 0.05 mg/ml cholesterol, but there was no difference in the total PI(4,5)P_2 levels in the DRM. One explanation for observing a statistically significant increase in motility after treatment with the same concentration of cholesterol could be that motility is very sensitive to the submembrane position of PI(4,5)P_2 and even small increases in PI(4,5)P_2 in rafts can cause enhanced cell movement. Our observation that an increase in PI(4,5)P_2 in rafts and uroids is accompanied by increases in intracellular Ca^{2+} and motility is suggestive of raft-mediated signaling from PI(4,5)P_2. However, it remains to be seen if the effect of PI(4,5)P_2 on motility after exposure to cholesterol is direct or indirect. One way to test this would be to generate *E. histolytica* cells with reduced levels of enzymes that are responsible for PI(4,5)P_2 synthesis (e.g., by knockdown of PIP5K). In *E. histolytica*, these enzymes have not yet been definitively identified. Genome data suggest the existence of a putative *E. histolytica* PIP5K gene (GI:67481998), and a BLASTP analysis of the translated protein sequence revealed a 41% identity to human phosphatidylinositol-4-phosphate-5-kinase type 1β. However, it remains to be seen if this an authentic PI(4,5)P_2-synthesizing enzyme.

Cholesterol is a major constituent of the eukaryotic plasma membrane (61). Aley et al. (62) demonstrated that *E. histolytica* possesses a high plasma membrane content of cholesterol, with a cholesterol/phospholipid molar ratio of 0.87. There are intriguing correlations between *E. histolytica* virulence and cholesterol. For example, cholesterol has been shown to enhance virulence functions in *E. histolytica*, including adhesion (27, 28), expression of ConA binding sites on the cell surface (63), and hemolytic and erythrophagocytic activity (64). Several studies have demonstrated that it is possible to revive the virulence of avirulent *E. histolytica* strains by supplying medium with cholesterol (65, 66) or by repeated passage through liver, which is a major site of cholesterol synthesis (67). Finally, long-term culture of *E. histolytica* in medium supplemented with liposomes carrying phosphatidylcholine and cholesterol can preserve virulence (63). Our study is the first to show that cholesterol impacts the submembrane and subcellular localization of PI(4,5)P_2 in *E. histolytica* and could be key to unraveling the molecular mechanism by which cholesterol regulates virulence.

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