

RESEARCH ARTICLE | *Molecular Pathways in Cell Signaling*

Integration of TRPC6 and NADPH oxidase activation in lysophosphatidylcholine-induced TRPC5 externalization

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Integration of TRPC6 and NADPH oxidase activation in lysophosphatidylcholine-induced TRPC5 externalization. *Am J Physiol Cell Physiol* 313: C541–C555, 2017. First published August 23, 2017; doi:10.1152/ajpcell.00028.2017.—Lipid oxidation products, including lysophosphatidylcholine (lysoPC), activate canonical transient receptor potential 6 (TRPC6) channels, and the subsequent increase in intracellular Ca²⁺ leads to TRPC5 activation. The goal of this study is to elucidate the steps in the pathway between TRPC6 activation and TRPC5 externalization. Following TRPC6 activation by lysoPC, extracellular regulated kinase (ERK) is phosphorylated. This leads to phosphorylation of p47^{phox} and subsequent NADPH oxidase activation with increased production of reactive oxygen species. ERK activation requires TRPC6 opening and influx of Ca²⁺ as evidenced by the failure of lysoPC to induce ERK phosphorylation in TRPC6^{-/-} endothelial cells. ERK siRNA blocks the lysoPC-induced activation of NADPH oxidase, demonstrating that ERK activation is upstream of NADPH oxidase. The reactive oxygen species produced by NADPH oxidase promote myosin light chain kinase (MLCK) activation with phosphorylation of MLC and TRPC5 externalization. Downregulation of ERK, NADPH oxidase, or MLCK with the relevant siRNA prevents TRPC5 externalization. Blocking MLCK activation prevents the prolonged rise in intracellular calcium levels and preserves endothelial migration in the presence of lysoPC.

TRPC6; TRPC5; NADPH oxidase; calcium; MLCK; endothelial

ENDOTHELIAL CELL (EC) migration is essential for the healing of arterial injuries such as those that accompany angioplasty. Rapid reendothelialization of an area of EC denudation limits the proliferation of smooth muscle cells and the subsequent development of intimal hyperplasia (2, 15, 23, 47, 48). Unfortunately, EC migration in vitro is inhibited by oxidized LDL and lipid oxidation products, including lysophosphatidylcholine (lysoPC) (25, 26), that are abundant in plasma and lesions of patients with atherosclerosis (9, 19, 31). Hypercholesterolemia inhibits reendothelialization of arterial injuries in vivo, and the delay in healing correlates with the elevation of plasma lysoPC (32, 34).

Normal cell migration requires exquisite control of calcium transients to allow detachment of focal adhesions, cytoskeletal reorganization to move the cell forward, and formation of new

attachments to the substratum. EC monolayer disruption is followed by influx of Ca²⁺ from the extracellular milieu and a transient rise in intracellular Ca²⁺ concentration ([Ca²⁺]_i) that is required for detachment of focal adhesions and initiation of movement (41). Following the initial rise in [Ca²⁺]_i, caveolae and the machinery for calcium wave initiation relocate to the trailing edge of the cell (20), allowing new focal adhesions to assemble in the leading portion of the cell. A generalized, sustained increase in [Ca²⁺]_i, such as that induced by lipid oxidation products, disrupts the time- and site-specific changes in focal adhesions and cytoskeleton that are required for cell movement and thus inhibits EC migration required to repair monolayer disruption (6).

We have shown that oxidized LDL and lysoPC, the primary lysophospholipid in oxidized LDL, inhibit EC migration, in part, by causing Ca²⁺ influx through canonical transient receptor potential (TRPC) channels (5). Specifically, lysoPC activates TRPC6 and the subsequent brief increase in [Ca²⁺]_i leads to externalization of TRPC5 and a prolonged increase in [Ca²⁺]_i. The increase in [Ca²⁺]_i activates calpains causing breakdown of cytoskeletal proteins and inhibition of EC migration (6). The importance of this pathway in vivo is demonstrated by the finding that a high-cholesterol diet dramatically inhibits endothelial healing of a carotid injury in wild-type (WT) mice but has no effect in TRPC6-deficient (*TRPC6*^{-/-}) mice (33). The mechanism of TRPC channel activation by lipid oxidation products is not completely elucidated.

Ca²⁺-dependent regulation of Ca²⁺-permeable cation channels is common, and all TRPC proteins have binding regions for calmodulin (CaM), a ubiquitous Ca²⁺-binding protein that can regulate TRPC activation (40). We have shown that lysoPC induces phosphorylation of CaM at Tyr⁹⁹, and this phosphorylation is required for the dissociation of CaM and TRPC6 and externalization of TRPC6 in the cell membrane (8). Interestingly, lysoPC does not cause dissociation of CaM and TRPC5, but lysoPC-induced TRPC5 externalization does require TRPC6 activation and influx of calcium (5, 8). The mechanism by which TRPC6 activation leads to TRPC5 externalization is the topic of this study.

TRPC5 channels can be activated by a multiplicity of signals including G protein-coupled receptor activation, extracellular gadolinium or Ca²⁺, modest elevation of [Ca²⁺]_i, or depletion of intracellular Ca²⁺ stores (50), but TRPC5 channel activation is inhibited by high [Ca²⁺]_i (28). We have shown that lysoPC-induced activation of TRPC5 requires the presence of TRPC6

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channels and Ca^{2+} in the medium (5). To determine the mechanism by which an increase in $[\text{Ca}^{2+}]_i$ through TRPC6 channels leads to activation of TRPC5, we assessed the contribution of extracellular regulated kinase (ERK) activation, activation of NADPH oxidase, increased reactive oxygen species (ROS) production, and myosin light chain kinase (MLCK) activation in the externalization of TRPC5. Determination of the mechanism by which lysoPC activates TRPC5 and identifying components of the activation cascade from TRPC6 to TRPC5 provide opportunities to interrupt these events, restore normal endothelial migration, and promote healing after interventions in atherosclerotic arteries.

MATERIALS AND METHODS

EC isolation and culture. Bovine aortic ECs were isolated from fresh adult bovine aortas and cultured as previously described (6). ECs in *passage 4 to 10* were used for experiments. Under a protocol approved by the Cleveland Clinic Institutional Animal Care and Use Committee, mouse aortic ECs (MAECs) were harvested from 129Sv/C57BL/6J WT, *TRPC6*^{-/-}, or *TRPC5*^{-/-} mice (14, 30), as described previously (5). The genetic deletion of the specific TRPC channel was confirmed by PCR using DNA extracted from tail or toe biopsies. MAECs in *passage 3 or 4* were used for experiments.

Immunoprecipitation and immunoblot analysis of target protein. Confluent ECs were used after being made quiescent in serum-free Dulbecco's modified Eagle's medium for 3 h before experimental conditions were initiated. Immunoprecipitation of target protein was performed as described previously (7).

Immunoblot analysis was performed after proteins (60–80 μg /lane) were loaded in 4–20% gradient SDS-PAGE as previously described (5, 7) and detected by antibodies specific for TRPC6 (1:200; catalog no. 600-401-FQ5; Rockland, St. Louis, MO), TRPC5 (1:250; catalog no. 73-104; UC Davis/NIH NeuroMab Facility), ERK1/2 (1:1,000; catalog no. 4695; Cell Signaling Technology, Danvers, MA), phospho-ERK1/2 (Thr²⁰²/Tyr²⁰⁴; 1:1,000; catalog no. SC-16982; Santa Cruz Biotechnology, Santa Cruz, CA), p47^{phox} (1:500; catalog no. 43125; Cell Signaling Technology), phospho-p47^{phox} (Ser³⁴⁵; 1:200; catalog no. A8391; Assay Biotechnology, Sunnyvale, CA), MLCK (1:500; catalog no. MABT194; Millipore), phospho-MLCK (Tyr⁴⁷¹; 1:500; catalog no. SC-17183; Santa Cruz Biotechnology), MLC (1:250; catalog no. M4401; Sigma), and phospho-MLC (Thr¹⁸/Ser¹⁹; 1:500; catalog no. 3474; Cell Signaling Technology). After being washed, the horseradish peroxidase-conjugated secondary antibody was applied and the signal developed using a chemiluminescent reagent (Perkin Elmer-Cetus, Boston, MA) for 1 min, and the image was acquired on HyBlot CL autoradiography film (Denville Scientific). The blot was then stripped and reprobed using an anti-actin antibody (1:2,000; catalog no. MA1-744; Invitrogen). For im-

munoblot analysis of phosphorylated proteins, samples were divided into two aliquots, one used for the phosphorylated protein and the other used for total ERK, p47^{phox}, MLCK, or MLC, as indicated. Protein band density on the HyBlot CL autoradiography film was quantitated using ImageJ software (NIH, Bethesda, MD). The film was then scanned with an HP Scanjet G4050 using HP Photosmart Premier Software.

Detection of TRPC externalization by biotinylation assay. Externalization of TRPC6 or TRPC5 was determined by biotinylation assay as previously described (5). Briefly, cell surface proteins were biotinylated and complexed with streptavidin-agarose beads, beads were collected, and precipitated proteins were released, resolved by SDS-PAGE, and identified by immunoblot analysis. Total TRPC6 or TRPC5 protein was determined by immunoblot analysis of an aliquot of cell lysate removed before incubation with streptavidin-agarose beads.

Overexpression of mutant CaM. The full-length human CaM cDNA (MGC-7) was obtained from ATCC. PCR-based site-directed mutagenesis was used to generate a cDNA for a mutant CaM in which Tyr⁹⁹ was replaced with Phe (Phe⁹⁹-CaM; ExonBio, San Diego, CA), as previously described (8). ECs at 60% confluence were transiently transfected with 2 μg of plasmids containing pcDNA3.1-myc-His-human CaM (WT-CaM) or pcDNA3.1-myc-His-Phe⁹⁹-CaM using Effectene (Qiagen, Chatsworth, CA) according to the manufacturer's protocol. The effectiveness of transfection was verified after 48 h by fluorescence microscopy and immunoblot analysis of CaM as previously described (8).

Downregulation of target protein by small interfering RNA. Using a transfection kit (RNAiFect; Qiagen), ECs at 80% confluence were transiently transfected with the small interfering (si)RNA duplex of CaM isoform1 (20 nM; Santa Cruz Biotechnology), ERK1 (20 nM; Santa Cruz Biotechnology), ERK2 (20 nM; Santa Cruz Biotechnology), p47^{phox} subunit of NADPH oxidase (20 nM; Santa Cruz Biotechnology), or MLCK (30 nM; Santa Cruz Biotechnology) for 24 h. Downregulation of endogenous target protein level was examined after 48 h by immunoblot analysis. A negative control siRNA (NsiRNA; 20–40 nM; Ambion, Austin, TX) without homology to any known gene sequence was used as a negative control.

Detection of ROS production. Intracellular ROS production was measured by oxidation of di(acetoxymethyl ester)(6-carboxy-2'-7'-dichlorodihydrofluorescein diacetate (H2DCF-DA; Life Technologies, Grand Island, NY). ECs at 70% confluence were incubated with H2DCF-DA (10 μM) for 20 min and washed twice before addition of lysoPC for 15 min. Fluorescence was monitored using Leica fluorescence microscope (Heidelberg, Germany) with FITC filter. The fluorescence intensity was quantitated using NIH ImageJ software.

Measurement of $[\text{Ca}^{2+}]_i$. $[\text{Ca}^{2+}]_i$ was measured using the Ca^{2+} -binding fluorophore fura 2-AM (Life Technologies), as previously described (5, 6). ECs were cultured in 35-mm dishes designed for

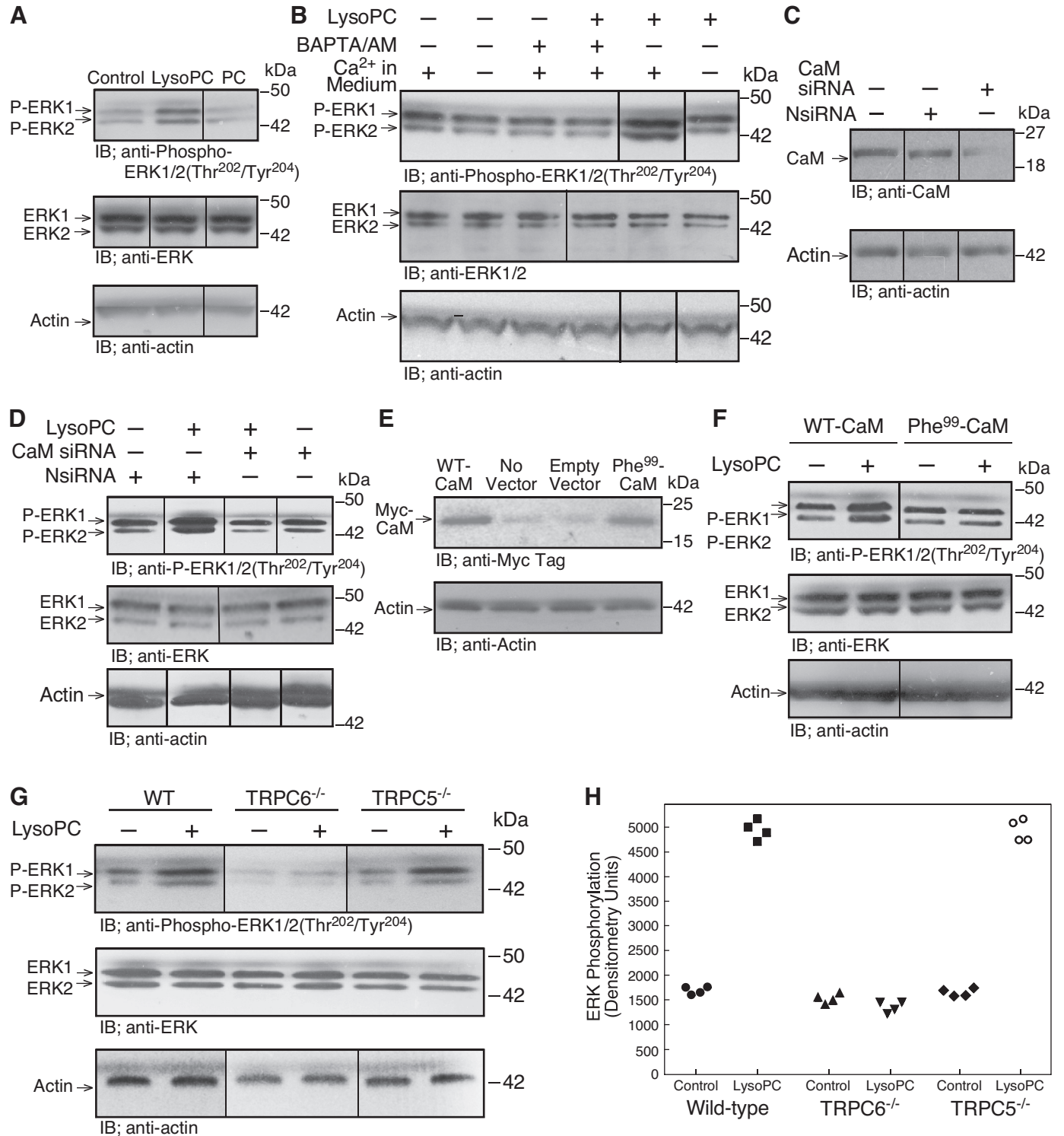
Fig. 1. LysoPC induces Ca^{2+} -, calmodulin (CaM)-, and TRPC6-dependent ERK phosphorylation. *A*: ECs were incubated with medium alone (Control), lysoPC (12.5 μM), or PC (12.5 μM) for 15 min. ERK and phospho-ERK1/2(Thr²⁰²/Tyr²⁰⁴) were detected by immunoblot (IB) analysis. Actin served as loading control (*n* = 4). *B*: ECs were incubated with calcium-free or calcium-containing KR buffer for 10 min or ECs were incubated with BAPTA/AM (25 μM) for 30 min before addition of lysoPC. Phospho-ERK1/2(Thr²⁰²/Tyr²⁰⁴) and total ERK1/2 were detected by immunoblot analysis. Actin served as loading control (*n* = 4). *C* and *D*: ECs were transiently transfected with control siRNA (NsiRNA) or CaM siRNA (20 nM) for 24 h. The siRNA was then removed. *C*: at 24 h after siRNA removal, cells were lysed and CaM was identified by immunoblot analysis. Actin served as loading control (*n* = 3). *D*: after siRNA removal, ECs were incubated with lysoPC (12.5 μM) for 15 min. Phospho-ERK1/2(Thr²⁰²/Tyr²⁰⁴) and total ERK1/2 were detected by immunoblot analysis. Actin served as loading control (*n* = 4). *E* and *F*: ECs were transiently transfected with pcDNA3.1-myc-His-WT-CaM or pcDNA3.1-myc-His-Phe⁹⁹-CaM for 24 h. *E*: overexpression of CaM was confirmed after 48 h by immunoblot analysis with anti-Myc Tag antibody. Actin served as loading control (*n* = 3). *F*: after transfection, ECs were incubated with lysoPC (12.5 μM) for 15 min. Phospho-ERK1/2(Thr²⁰²/Tyr²⁰⁴) and total ERK were detected by immunoblot analysis. Actin served as loading control (*n* = 4). *G* and *H*: WT, *TRPC6*^{-/-}, or *TRPC5*^{-/-} MAECs were incubated with lysoPC (10 μM) for min. *G*: phospho-ERK1/2(Thr²⁰²/Tyr²⁰⁴) and total ERK were detected by immunoblot analysis. Actin served as loading control (*n* = 3). *H*: densitometry measurements of phospho-ERK(Thr²⁰²/Tyr²⁰⁴) are represented in graphic form: ●, medium control with WT ECs; ■, lysoPC with WT ECs; ▲, medium control with *TRPC6*^{-/-} ECs; ▼, lysoPC with *TRPC6*^{-/-} ECs; ◆, medium control with *TRPC5*^{-/-} ECs; ○, lysoPC with *TRPC5*^{-/-} ECs. (In A–G, lines indicate lanes rearranged from the same gel. All bands are from the same gel.)

fluorescence microscopy (Bioptechs, Butler, PA), incubated with fura 2-AM (1 μ M) for 30 min, and washed with Krebs-Ringer (KR) buffer to remove excess fura 2-AM. Fluorescence of a group of 10–12 cells inside the light path was continuously examined with an Olympus IX-70 inverted fluorescence microscope (Melville, NY). The relative change of $[Ca^{2+}]_i$ was estimated by the ratio of fura 2 excitation at wavelengths of 340 nm and 380 nm (340/380 ratio).

EC migration assay. ECs were made quiescent in serum-free Dulbecco's modified Eagle's medium for 8–12 h before the migration

assay. EC migration was assessed by the razor-scrape assay as described in our previous study (6). Migrating cells were quantitated using NIH Image 1.63 software by an observer blinded to the experimental conditions.

Statistics. Results are presented as means \pm SD. Experiments were performed in triplicate with ECs isolated from at least three different animals. Statistical analysis was performed using Student's *t*-test or ANOVA, and differences between means were considered statistically significant at $P < 0.05$.



RESULTS

LysoPC induced Ca²⁺-CaM dependent ERK phosphorylation. Previously, we showed that lysoPC (1-palmitol-2-hydroxi-*sn*-glycerol-3-phosphocholine; Avanti Polar Lipids, Alabaster, AL) increased [Ca²⁺]_i through a unique TRPC6 channel to TRPC5 channel activation cascade (5). An increase in [Ca²⁺]_i through TRPC6 channels can lead to ERK activation (18), and ERK can activate MLCK (22), which is essential in TRPC5 externalization (21, 37). Therefore, a possible role of ERK in the lysoPC-induced TRPC activation cascade was studied. In ECs incubated with lysoPC (12.5 μM), but not phosphatidylcholine (PC; 12.5 μM), for 15 min, ERK1/2 phosphorylation at Thr²⁰²/Tyr²⁰⁴ increased to 280 ± 30% of control ($n = 4$; $P < 0.01$; Fig. 1A). Total ERK levels did not change.

To determine if lysoPC-induced ERK phosphorylation was Ca²⁺ dependent, ECs were incubated in Ca²⁺-free or Ca²⁺-containing KR buffer or with the intracellular Ca²⁺-chelator BAPTA/AM (25 μM) before the addition of lysoPC (12.5 μM). LysoPC did not induce ERK1/2 phosphorylation in the absence of extracellular Ca²⁺ or presence of BAPTA/AM, suggesting that ERK1/2 phosphorylation was Ca²⁺ dependent ($n = 4$; $P < 0.01$ compared with incubation with lysoPC in Ca²⁺-containing KR buffer; Fig. 1B).

ERK activation can be regulated by CaM (1), and lysoPC induces CaM phosphorylation at Tyr⁹⁹ that is required for TRPC6 externalization (8). Transient transfection of ECs with CaM siRNA significantly reduced the level of CaM (Fig. 1C) and the lysoPC-induced ERK1/2 phosphorylation [$n = 4$; $P < 0.01$, compared with ECs transiently transfected with negative control siRNA (NsiRNA) that has no homology to a known gene sequence; Fig. 1D]. When ECs were transiently transfected with WT-CaM or mutant CaM in which the Tyr⁹⁹ site was replaced with Phe (Phe⁹⁹-CaM) that cannot be phosphorylated (Fig. 1E), lysoPC increased ERK1/2 phosphorylation in ECs transfected with WT-CaM but not in ECs overexpressing Phe⁹⁹-CaM ($n = 4$; $P < 0.01$, compared with WT-CaM; Fig. 1F). Taken together, these findings suggested that lysoPC-induced ERK activation in ECs was Ca²⁺-CaM dependent and required CaM phosphorylation at Tyr⁹⁹.

ERK activation by lysoPC was downstream of TRPC6 externalization but not TRPC5 externalization. CaM phosphorylation is required for TRPC6 externalization (8); therefore, the relationship between lysoPC-induced ERK activation and TRPC activation was explored using *TRPC6*^{-/-} and *TRPC5*^{-/-} MAECs. The basal level of ERK1/2 phosphorylation was similar in WT, *TRPC6*^{-/-}, and *TRPC5*^{-/-} MAECs under control conditions (Fig. 1G). Incubation with lysoPC (10 μM) for 15 min caused an increase in ERK1/2 phosphorylation to 301 ± 10% of control in WT MAECs ($n = 3$; $P < 0.01$; Fig. 1, G and H), but no change in ERK1/2 phosphorylation in *TRPC6*^{-/-} MAECs ($n = 3$; Fig. 1, G and H). Similar to WT MAECs, lysoPC induced an increase in ERK1/2 phosphorylation to 300 ± 10% of control in *TRPC5*^{-/-} MAECs ($n = 3$; $P < 0.03$; Fig. 1, G and H). These data indicated that lysoPC-induced ERK activation required TRPC6 but not TRPC5, suggesting that ERK activation is downstream from TRPC6 but upstream from TRPC5 activation.

To further assess the role of ERK in TRPC externalization, ERK was downregulated with siRNA. Transient transfection of ECs with ERK siRNA resulted in a significant decrease in ERK

at 48 h ($n = 3$; $P < 0.01$, compared with NsiRNA; Fig. 2A). The effect of ERK downregulation on TRPC6 and TRPC5 externalization was assessed by biotinylation assay (Fig. 2, B–G). Transfection with ERK siRNA had no effect on lysoPC-induced TRPC6 externalization ($n = 4$; Fig. 2, D and E); however, TRPC5 externalization was significantly inhibited ($n = 4$; $P < 0.04$, compared with NsiRNA transfected ECs incubated with lysoPC; Fig. 2, F and G). These results suggested that lysoPC increased TRPC5 externalization through an ERK-dependent mechanism.

LysoPC induced ERK-dependent NADPH activation. LysoPC and ERK can activate NADPH oxidase (35, 43), and TRPC5 can be activated by ROS (27). Therefore, we explored the role of NADPH oxidase in lysoPC-induced TRPC5 externalization. We previously showed that lysoPC caused the p47^{phox} and p67^{phox} subunits of NADPH oxidase to translocate from the cytosol to the membrane (32), characteristic of NADPH oxidase activation. After EC incubation in lysoPC (12.5 μM), but not PC (12.5 μM), for 15 min, p47^{phox} phosphorylation at Ser³⁴⁵ increased to 340 ± 20% of control ($n = 4$; $P < 0.02$; Fig. 3A), again suggesting NADPH oxidase activation.

To determine if lysoPC-induced NADPH oxidase activation was Ca²⁺ dependent, ECs were incubated in Ca²⁺-free or Ca²⁺-containing KR buffer, or with BAPTA/AM (25 μM), before addition of lysoPC (12.5 μM). LysoPC did not induce p47^{phox} phosphorylation at Ser³⁴⁵ in absence of extracellular Ca²⁺ ($n = 4$; $P < 0.01$, compared with Ca²⁺-containing KR buffer; Fig. 3B) or in the presence of BAPTA/AM ($n = 4$; $P < 0.02$, compared with incubation with lysoPC in Ca²⁺-containing KR buffer; Fig. 3B). In addition, lysoPC-induced translocation of p47^{phox} to the membrane from the cytosol was reduced in the absence of Ca²⁺ ($n = 4$; Fig. 3C), suggesting that NADPH oxidase activation was Ca²⁺ dependent.

The role of ERK activation in NADPH oxidase activation was explored. LysoPC induced ERK phosphorylation, and this was accompanied by increased association of p47^{phox} and phospho-ERK (Fig. 3D; $n = 3$) as well as p67^{phox} and phospho-ERK (Fig. 3E; $n = 3$). To determine if NADPH oxidase activation by lysoPC required ERK, ERK was downregulated with siRNA for 24 h and then, lysoPC (12.5 μM) was added for 15 min. ERK downregulation caused a significant reduction in lysoPC-induced p47^{phox} phosphorylation ($n = 5$; $P < 0.02$, compared with NsiRNA; Fig. 3F), indicating that lysoPC-induced NADPH oxidase activation was ERK dependent. Conversely, downregulation of p47^{phox} did not alter lysoPC-induced ERK phosphorylation ($n = 4$; Fig. 3G). Similarly, downregulation of p22^{phox} ($n = 3$; Fig. 3H) did not alter lysoPC-induced ERK phosphorylation ($n = 3$; Fig. 3I).

To confirm that p47^{phox} phosphorylation induced by lysoPC required TRPC6 activation, p47^{phox} phosphorylation was assessed in *TRPC6*^{-/-} or *TRPC5*^{-/-} MAECs. LysoPC induced p47^{phox} phosphorylation in WT MAECs but not in *TRPC6*^{-/-} MAECs ($n = 4$; $P < 0.01$, compared with WT MAECs with lysoPC; Fig. 4, A and B). LysoPC-induced p47^{phox} phosphorylation in *TRPC5*^{-/-} MAECs was similar to that in WT MAECs ($n = 4$; Fig. 4, A and B). These results suggested that lysoPC-induced NADPH oxidase activation was TRPC6 dependent. To determine if TRPC6 was required for NADPH oxidase activation by lysoPC in human ECs, TRPC6 was downregulated using siRNA ($n = 3$; Fig. 4C). This blocked lysoPC-induced p47^{phox} phosphorylation ($n = 3$; Fig. 4D).

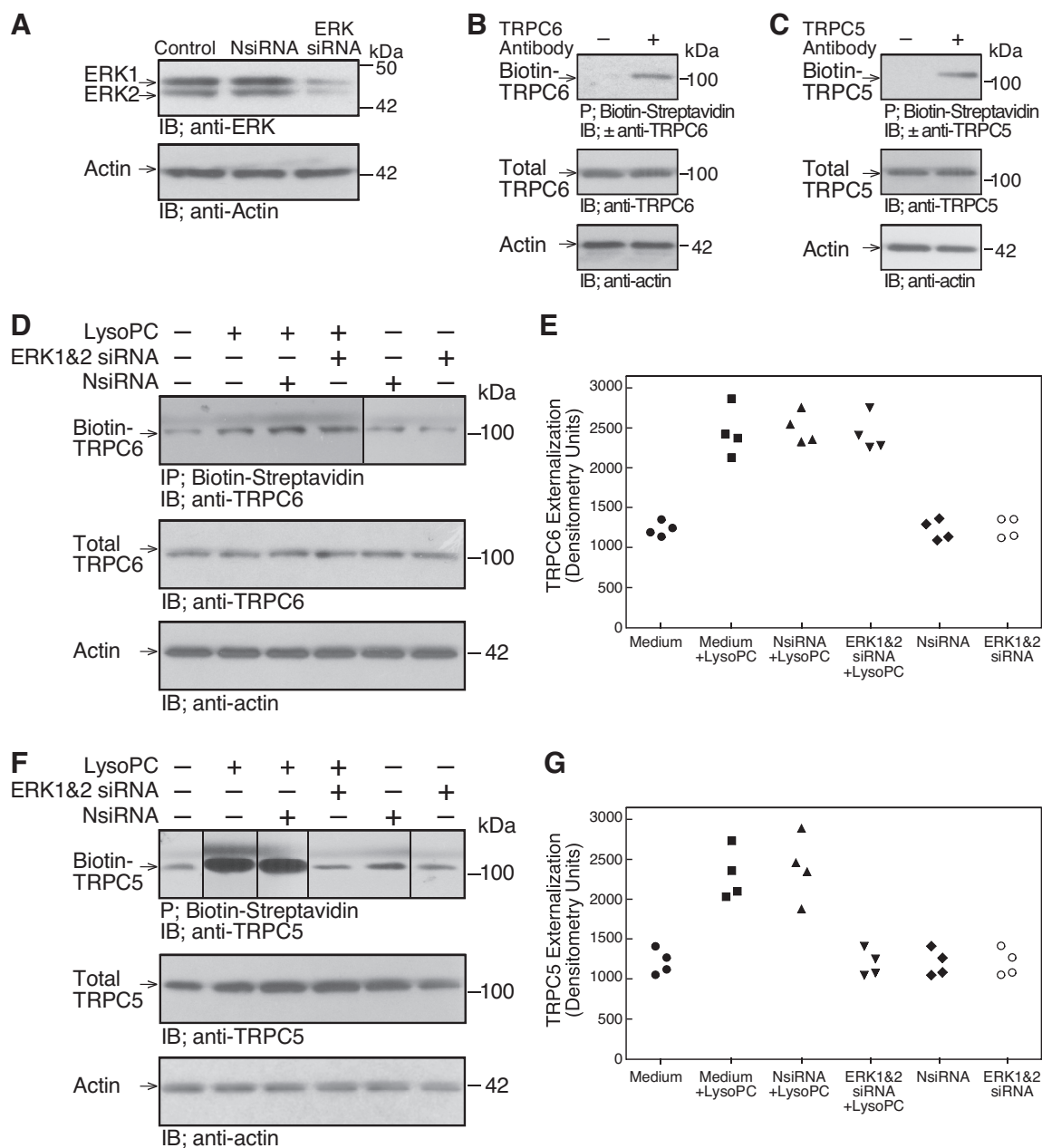


Fig. 2. LysoPC-induced ERK activation regulates TRPC5 externalization. *A*: ECs were transiently transfected with NsiRNA (40 nM) or ERK1 siRNA and ERK2 siRNA (20 nM of each) for 24 h. The siRNA was removed, and 24 h later the cells were lysed. ERK was detected by immunoblot analysis. Actin served as loading control ($n = 3$). *B* and *C*: TRPC externalization was determined by biotinylation of proteins on the cell surface, precipitation of biotin-streptavidin complexes, resolution of proteins by SDS-PAGE, and immunoblot analysis. To verify the specificity of the assay, controls included omission of the primary antibody for TRPC6 (*B*) or TRPC5 (*C*) but inclusion of the horseradish peroxidase-conjugated secondary antibody. Total TRPC6 or TRPC5 was determined by immunoblot analysis of an aliquot of the same sample ($n = 3$). Actin served as loading control ($n = 3$). *D–G*: ECs were transiently transfected with NsiRNA or ERK1 siRNA and ERK2 siRNA for 24 h before incubation with lysoPC (12.5 μ M) for 15 min. Externalization of TRPC6 (*D*) or TRPC5 (*F*) was detected by biotinylation assay and total TRPC6 (*D*) or TRPC5 (*F*) by immunoblot analysis. Actin served as loading control ($n = 4$). The increase in externalized TRPC6 (*E*) or TRPC5 (*G*) is represented in graphic form: ●, medium control; ■, medium + lysoPC; ▲, NsiRNA + lysoPC; ▼, ERK 1 and 2 siRNA + lysoPC; ◆, NsiRNA medium control; ○, ERK 1 and 2 siRNA medium control. (In *D* and *F*, lines indicate lanes rearranged from the same gel. All bands are from the same gel.)

Superoxide and NADPH oxidase activation were required for lysoPC-induced TRPC5 externalization. Because ROS can activate TRPC5 (27), the role of superoxide in lysoPC-induced TRPC6 and TRPC5 externalization was assessed by biotinylation assay. ECs were pretreated with superoxide dismutase (SOD; 50 units/ml; Sigma) or PEG-catalase (5,000 units/ml) for 30 min before incubation with lysoPC (12.5 μ M) for 15

min. LysoPC-induced TRPC6 externalization was not altered by either agent ($n = 3$; Fig. 5A). TRPC5 externalization, however, was blocked by SOD, suggesting that superoxide production was required for lysoPC-induced TRPC5 externalization ($n = 3$; Fig. 5B). Preincubation of ECs with PEG-catalase did not block lysoPC-induced TRPC5 externalization (Fig. 5B).

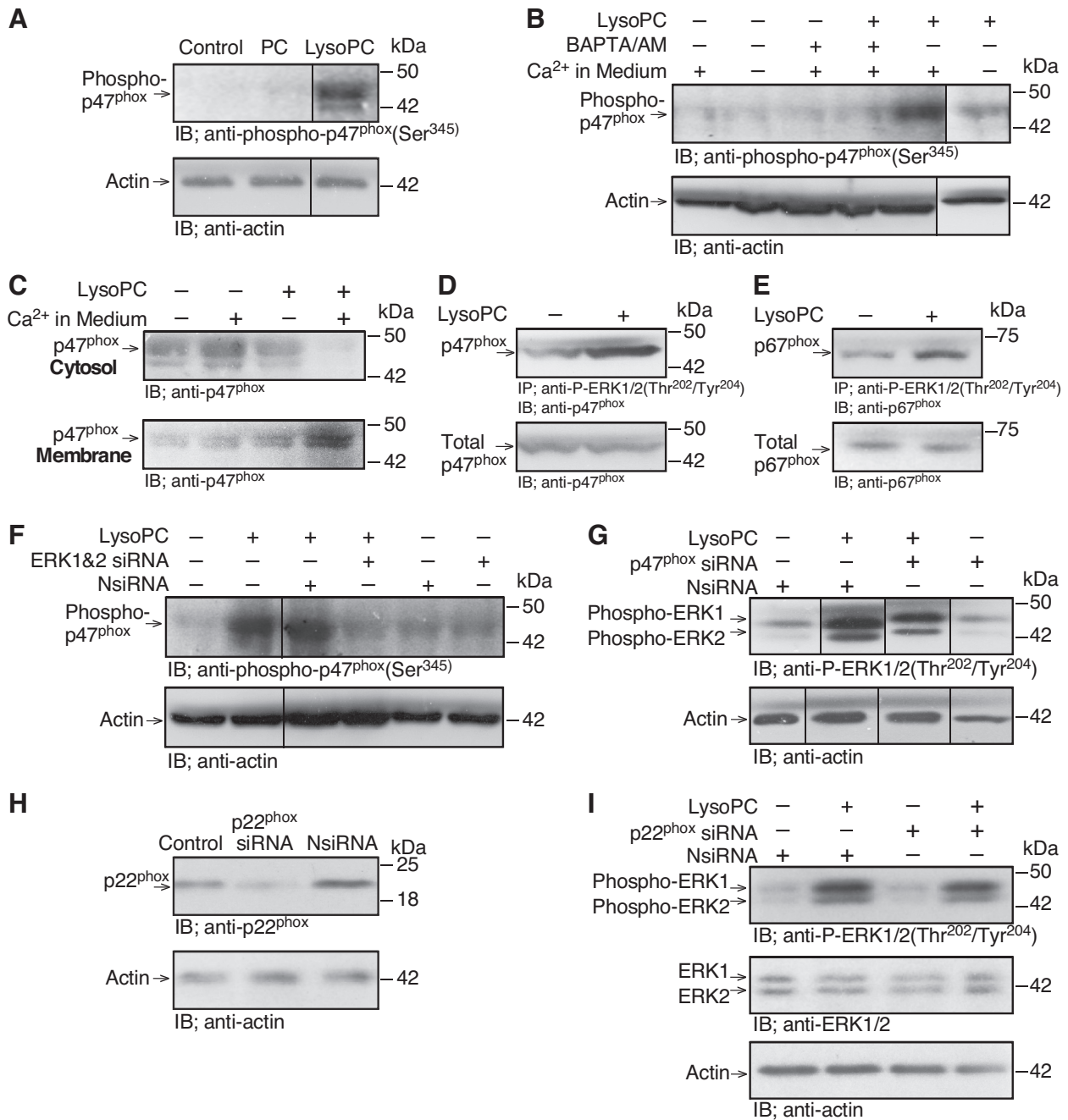


Fig. 3. LysoPC-induced NADPH oxidase activation is Ca²⁺- and ERK dependent. **A:** ECs were incubated with PC (12.5 μM) or lysoPC (12.5 μM) for 15 min. Phospho-p47^{phox}(Ser³⁴⁵) was detected by immunoblot analysis, and actin served as loading control (n = 4). **B:** ECs were incubated in calcium-free or calcium-containing KR buffer for 10 min or ECs were incubated with BAPTA/AM (25 μM) for 30 min before adding lysoPC. Phospho-p47^{phox}(Ser³⁴⁵) was detected by immunoblot analysis. Actin served as loading control (n = 4). **C:** ECs were incubated with calcium-free or calcium-containing KR buffer for 10 min before adding lysoPC. Cytosolic and membrane fraction proteins were separated, and p47^{phox} was detected by immunoblot analysis (n = 4). **D and E:** ECs were incubated with lysoPC (12.5 μM) for 15 min. Phospho-ERK1/2(Thr²⁰²/Tyr²⁰⁴) was immunoprecipitated, and associated p47^{phox} (**D**) or p67^{phox} (**E**) was detected by immunoblot analysis (n = 3). Total p47^{phox} or p67^{phox} was measured by immunoblot analysis of the lysate before immunoprecipitation (n = 3). **F:** ECs were transiently transfected with NsiRNA (40 nM) or ERK1 siRNA and ERK2 siRNA (20 nM of each) for 24 h before incubation with lysoPC. Phospho-p47^{phox}(Ser³⁴⁵) was detected by immunoblot analysis, and actin served as loading control (n = 5). **G:** ECs were transfected with NsiRNA or p47^{phox} siRNA (20 nM) for 24 h, then incubated with lysoPC (12.5 μM) for 15 min. ERK phosphorylation (n = 4) was detected by immunoblot analysis. Actin served as loading control (n = 4). **H and I:** ECs were transfected with NsiRNA or p22^{phox} siRNA (30 nM) for 24 h. The siRNA was then removed. **H:** At 24 h after siRNA removal, the cells were lysed, and p22^{phox} was identified by immunoblot analysis. Actin served as loading control (n = 3). **I:** After siRNA removal, ECs were incubated with lysoPC (12.5 μM) for 15 min. Phospho-ERK1/2(Thr²⁰²/Tyr²⁰⁴) and ERK1/2 were detected by immunoblot analysis. Actin served as loading control (n = 3). (In **A** and **B**, **F** and **G**, lines indicate lanes rearranged from the same gel. All bands are from the same gel.)

To verify the role NADPH oxidase activity in the lysoPC-induced superoxide production, p47^{phox} was downregulated with siRNA. In ECs transiently transfected with p47^{phox} siRNA for 48 h, p47^{phox} protein levels were significantly reduced ($n = 5$; $P < 0.01$, compared with NsiRNA; Fig. 5C). Next, ROS production was determined using H2DCF-DA. Addition of lysoPC (12.5 μ M) for 15 min induced an increase in ROS production to $780 \pm 40\%$ of control in NsiRNA-transfected ECs as detected by H2DCF-DA assay ($n = 5$; $P <$

0.01; Fig. 5, D and E). In ECs transiently transfected with p47^{phox} siRNA, incubation with lysoPC caused an increase in ROS production to only $201 \pm 40\%$ of control ($n = 5$; $P < 0.02$ compared with NsiRNA-transfected ECs incubated with lysoPC; Fig. 5, D and E). Thus downregulation of p47^{phox} decreased the lysoPC-induced ROS production by $\sim 75\%$.

Next, the role of NADPH oxidase in TRPC6 and TRPC5 externalization was assessed. In ECs transiently transfected with p47^{phox} siRNA, incubation with lysoPC induced TRPC6 externalization ($n = 4$; Fig. 5F), but TRPC5 externalization was blocked ($n = 5$; $P < 0.02$, compared with NsiRNA with lysoPC; Fig. 5G). In ECs incubated for 30 min with TNF- α (500 U/ml), an NADPH oxidase activator (17), TRPC6 was not externalized ($n = 3$; Fig. 5H), but TRPC5 externalization was increased to $202 \pm 40\%$ of control ($n = 3$; $P < 0.01$; Fig. 5I). These results suggested that NADPH oxidase activation could promote TRPC5 externalization but not TRPC6 externalization.

LysoPC-induced MLCK activation required Ca²⁺ influx and TRPC6 channels. ROS can activate MLCK (51), a Ca²⁺-CaM-dependent kinase that phosphorylates MLC, and MLC can modulate the insertion of TRPC5 containing vesicles into the plasma membrane, resulting in activation of TRPC5 (21, 37). We assessed MLCK phosphorylation at Tyr⁴⁷¹ as this can contribute to increased enzymatic activity (3). In ECs incubated with lysoPC (12.5 μ M), but not PC (12.5 μ M), for 15 min, MLCK phosphorylation at Tyr⁴⁷¹ increased to $230 \pm 40\%$ of medium alone ($n = 3$; $P < 0.01$; Fig. 6A). LysoPC-induced MLCK phosphorylation was inhibited in the absence of extracellular Ca²⁺ ($n = 4$; $P < 0.03$, compared with Ca²⁺-containing KR buffer; Fig. 6B) or the presence of BAPTA/AM ($n = 4$; $P < 0.01$, compared with Ca²⁺-containing KR buffer; Fig. 6B), suggesting that lysoPC-induced MLCK phosphorylation was Ca²⁺ dependent.

To determine if MLCK activation was dependent on TRPC activation, we assessed MLCK phosphorylation in WT, TRPC6^{-/-}, or TRPC5^{-/-} MAECs. LysoPC induced a robust increase in MLCK phosphorylation to $220 \pm 30\%$ of control in WT MAECs ($n = 4$; $P < 0.02$) and an increase in MLCK phosphorylation to $220 \pm 26\%$ of control in TRPC5^{-/-} MAECs ($n = 4$, $P < 0.02$; Fig. 6, C and D). No increase was noted when TRPC6^{-/-} MAECs were incubated with lysoPC ($n = 4$; $P = 0.7$, compared with WT control; Fig. 6, C and D), confirming that lysoPC-induced MLCK phosphorylation was TRPC6 dependent. The MLCK phosphorylation was accom-

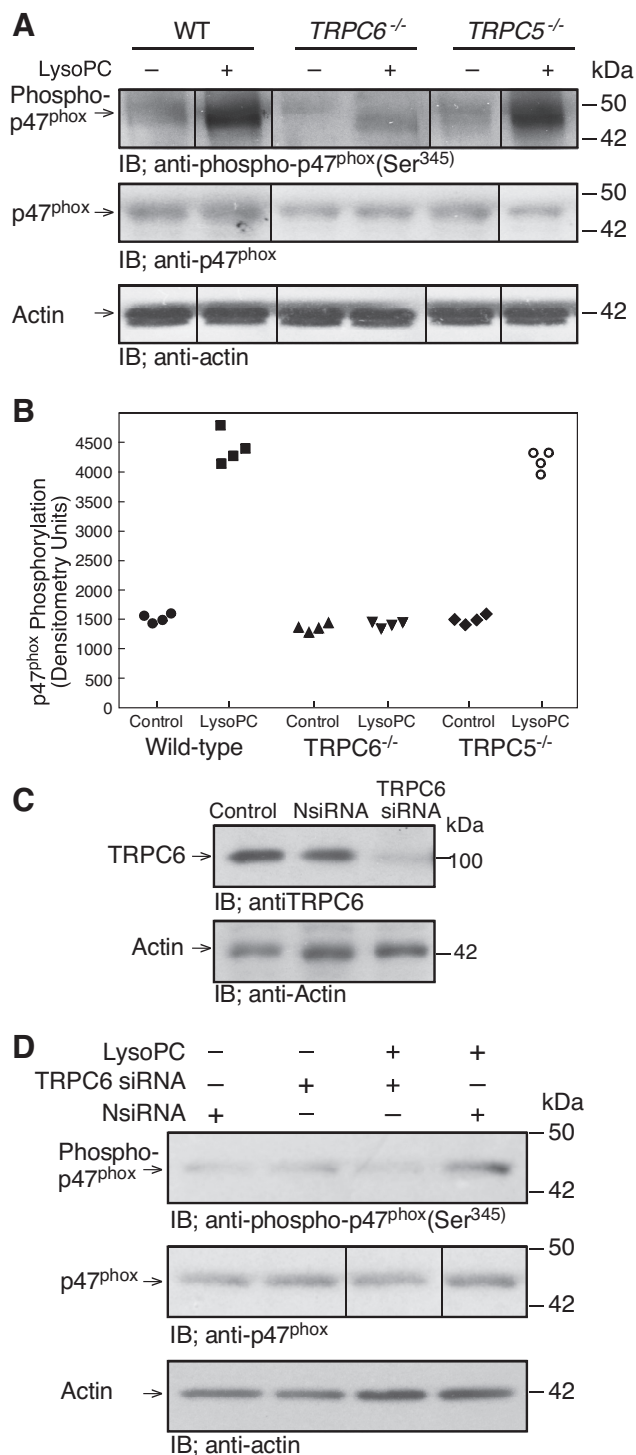
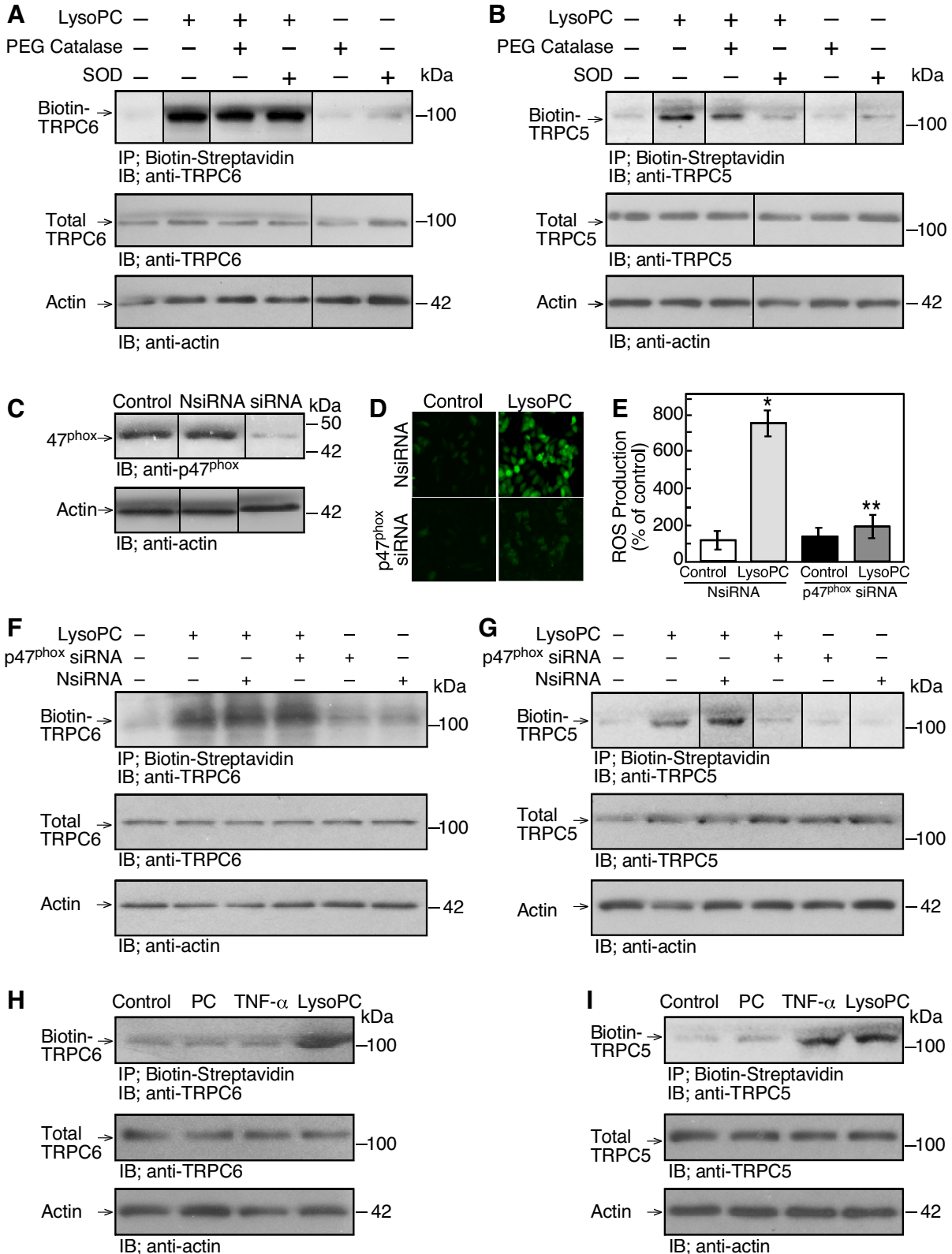


Fig. 4. LysoPC-induced NADPH oxidase activation requires TRPC6. A and B: WT, TRPC6^{-/-}, or TRPC5^{-/-} MAECs were incubated with lysoPC (10 μ M) for 15 min. A: phospho-p47^{phox}(Ser³⁴⁵) and total p47^{phox} were detected by immunoblot analysis. Actin served as loading control ($n = 4$). B: densitometry measurements of and phospho-p47^{phox}(Ser³⁴⁵) are represented in graphic form: ●, medium control with WT ECs; ■, lysoPC with WT ECs; ▲, medium control with TRPC6^{-/-} ECs; ▼, lysoPC with TRPC6^{-/-} ECs; ◆, medium control with TRPC5^{-/-} ECs; ○, lysoPC with TRPC5^{-/-} ECs. C and D: human ECs (EA.hy926) were transiently transfected with NsiRNA (20 nM) or TRPC6 siRNA (20 nM) for 24 h. siRNA was removed. C: at 24 h after siRNA removal, cells were lysed and TRPC6 was detected by immunoblot analysis. Actin served as loading control ($n = 3$). D: after siRNA removal, EA.hy926 cells were incubated with lysoPC (12.5 μ M) for 15 min. Phosphorylation of p47^{phox} and total p47^{phox} were detected by immunoblot analysis. Actin served as loading control ($n = 3$). (In A and D, lines indicate lanes rearranged from the same gel. All bands are from the same gel.)

panied by an increase in MLC phosphorylation. Incubation with lysoPC increased MLC phosphorylation to $230 \pm 40\%$ of control in WT MAECs ($n = 4$; $P < 0.02$; Fig. 6, E and F). LysoPC also increased MLC phosphorylation to $220 \pm 30\%$ of

control in *TRPC5*^{-/-} MAECs ($n = 4$; $P < 0.02$; Fig. 6, E and F). No lysoPC-induced increase in MLC phosphorylation was noted in *TRPC6*^{-/-} MAECs ($n = 4$; $P = 0.8$, compared with *TRPC6*^{-/-} control; Fig. 6, E and F). The endogenous levels of



MLC were not changed in the presence of lysoPC ($n = 3$; Fig. 6E). These results supported the TRPC6 dependence of MLCK activation by lysoPC.

ERK and NADPH oxidase activation were required for lysoPC-induced MLCK activation and MLC phosphorylation. The role of ERK and NADPH oxidase in MLCK phosphorylation was assessed. In ECs incubated with the NADPH oxidase activator TNF- α (500 U/ml) for 30 min, MLCK phosphorylation was increased significantly ($n = 3$; $P < 0.03$, compared with control; Fig. 7A). The roles of ERK and NADPH oxidase in MLCK phosphorylation were confirmed by downregulating ERK or p47^{phox}. ERK siRNA significantly reduced lysoPC-induced MLCK phosphorylation ($n = 4$; $P < 0.01$, compared with NsiRNA; Fig. 7B). Similarly, p47^{phox} siRNA inhibited lysoPC-induced MLCK phosphorylation ($n = 4$; $P < 0.02$, compared with NsiRNA; Fig. 7B), suggesting the involvement of both ERK and NADPH oxidase in lysoPC-induced MLCK phosphorylation. The lysoPC-induced MLCK phosphorylation was accompanied by an increase in MLC phosphorylation. In the medium control, lysoPC increased MLC phosphorylation to $240 \pm 20\%$ of control ($n = 3$; $P < 0.01$; Fig. 7C). In ECs transfected with NsiRNA, lysoPC also increased MLC phosphorylation to $240 \pm 20\%$ of control ($n = 3$; $P < 0.01$; Fig. 7C). Downregulation of ERK using siRNA blocked lysoPC-induced MLC phosphorylation ($n = 3$; $P < 0.02$, compared with NsiRNA with lysoPC; $P < 0.01$, compared with medium control; Fig. 7C). Similarly, downregulation of p47^{phox} using siRNA significantly reduced lysoPC-induced MLC phosphorylation ($n = 3$; $P < 0.01$, compared with NsiRNA with lysoPC; Fig. 7D). These results suggested that lysoPC-induced MLCK phosphorylation and MLC phosphorylation were dependent on the activation of ERK and NADPH oxidase.

MLCK activation by lysoPC required superoxide. The results with p47^{phox} siRNA suggested that superoxide production is required for lysoPC-induced MLCK activation. To confirm the role of ROS, MLCK phosphorylation was measured in the presence of SOD. ECs were preincubated with SOD for 30 min and then lysoPC (12.5 μ M) was added for 15 min. Incubation with lysoPC increased MLCK phosphorylation to $260 \pm 20\%$ of control ($n = 3$; $P < 0.01$; Fig. 7E), but after SOD pretreatment lysoPC did not cause a significant increase ($n = 3$; $P = 0.8$, compared with SOD control; Fig. 7E). Similarly, lysoPC increased MLC phosphorylation to $260 \pm 10\%$ of medium control ($n = 3$; $P < 0.01$; Fig. 7F), but after SOD pretreatment lysoPC did not cause a significant increase ($n = 3$; $P = 0.8$, compared with SOD control; Fig. 7F). Endogenous levels of

MLCK and MLC were not changed by SOD pretreatment ($n = 3$; Fig. 7, E and F). These results suggested that lysoPC-induced MLCK activation required superoxide.

LysoPC-induced TRPC5 externalization required MLCK phosphorylation. The role of MLCK in TRPC5 externalization was analyzed by inhibiting or downregulating MLCK. ECs were preincubated with ML-9 (10 μ M), a relatively selective inhibitor that impairs MLCK phosphorylation by competing for the ATP binding site. After 30 min, lysoPC (12.5 μ M) was added for 15 min. ML-9 had no effect on total TRPC6 or TRPC5 level and did not inhibit the lysoPC-induced TRPC6 translocation to the cell membrane suggesting that TRPC6 activation was MLCK independent ($n = 4$; Fig. 8A). However, ML-9 blocked lysoPC-induced TRPC5 translocation ($n = 4$; $P < 0.02$, compared with lysoPC control; Fig. 8B). In ECs transiently transfected with MLCK siRNA for 48 h, MLCK was downregulated significantly ($n = 4$; $P < 0.02$, compared with NsiRNA; Fig. 8C). When MLCK was downregulated, lysoPC-induced TRPC6 externalization was unchanged ($n = 4$, Fig. 8D), but TRPC5 externalization was markedly diminished ($n = 4$; $P < 0.02$, compared with NsiRNA with lysoPC; Fig. 8E). Similarly, when MLCK was downregulated with siRNA, incubation with TNF- α (500 U/ml) for 30 min did not increase TRPC5 externalization ($n = 4$; $P = 0.8$, compared with MLCK siRNA-transfected ECs without TNF- α ; Fig. 8E). These results suggested that NADPH oxidase-mediated TRPC5 externalization, including that by lysoPC, was dependent on MLCK.

MLCK inhibition reduced lysoPC-induced $[Ca^{2+}]_i$ rise and inhibition of EC migration. To determine if inhibition of MLCK blocked the lysoPC-mediated increase of $[Ca^{2+}]_i$, ECs were loaded with fura 2-AM for 30 min. After baseline $[Ca^{2+}]_i$ readings were obtained, lysoPC (12.5 μ M) was added and the increase of $[Ca^{2+}]_i$ was assessed. LysoPC was removed and cells were washed. Then, ECs were incubated with ML-9 (10 μ M) for 20 min and lysoPC again added. ML-9 did not alter the basal $[Ca^{2+}]_i$ readings nor inhibit the initial transient increase in $[Ca^{2+}]_i$ induced by lysoPC, presumably through TRPC6 channels but blocked the long plateau phase ($n = 4$; $P < 0.02$, compared with lysoPC control; Fig. 8F) that was attributed to TRPC5 activation (5).

To determine the role of MLCK in lysoPC-induced inhibition of EC migration, ECs were preincubated with ML-9 for 30 min. Basal migration of ECs with or without ML-9 was similar. LysoPC inhibited 24 h EC migration by 63%, but after pretreatment with ML-9, lysoPC inhibited EC migration by only 42% ($n = 5$; $P < 0.001$, compared with lysoPC without ML-9; Fig. 8G). Since ML-9 is only relatively specific for

Fig. 5. NADPH oxidase activation is required for TRPC5 externalization. A and B: ECs were made serum-free for 3 h, then pretreated with superoxide dismutase (SOD; 50 units/ml) or PEG catalase (5,000 units/ml) for 30 min before incubation with lysoPC (12.5 μ M) for 15 min. TRPC6 or TRPC5 externalization was detected by biotinylation assay and total TRPC6 or TRPC5 was determined by immunoblot analysis of an aliquot of the same sample ($n = 3$). Actin served as loading control ($n = 3$). C: ECs were transiently transfected with NsiRNA or p47^{phox} siRNA (20 nM) for 24 h. The siRNA was removed, and 24 h later cells were lysed. Total p47^{phox} was detected by immunoblot analysis. Actin served as loading control ($n = 5$). D and E: ECs were transiently transfected with NsiRNA or p47^{phox} siRNA for 24 h and then incubated with H2DCF-DA for 20 min before incubation with lysoPC ($n = 5$). D: ROS production was detected by DCF assay using fluorescence microscopy. E: fluorescence intensity was quantitated using image analysis software and graphed. Results are represented as means \pm SD ($n = 5$; * $P < 0.01$, compared with NsiRNA-transfected EC control; ** $P < 0.02$, compared with NsiRNA-transfected ECs incubated with lysoPC). F and G: ECs were transiently transfected with NsiRNA or p47^{phox} siRNA for 24 h before incubation with lysoPC. Externalization of TRPC6 or TRPC5 was detected by biotinylation assay and total TRPC6 or TRPC5 by immunoblot analysis. Actin served as loading control (F: $n = 4$; G: $n = 5$). H and I: ECs were incubated with PC (12.5 μ M) for 15 min, TNF- α (500 U/ml) for 30 min, or lysoPC (12.5 μ M) for 15 min. Externalization of TRPC6 or TRPC5 was detected by biotinylation assay and total TRPC6 or TRPC5 by immunoblot analysis. Actin served as loading control ($n = 3$). (In A–C and G, lines indicate lanes rearranged from the same gel. All bands are from the same gel.)

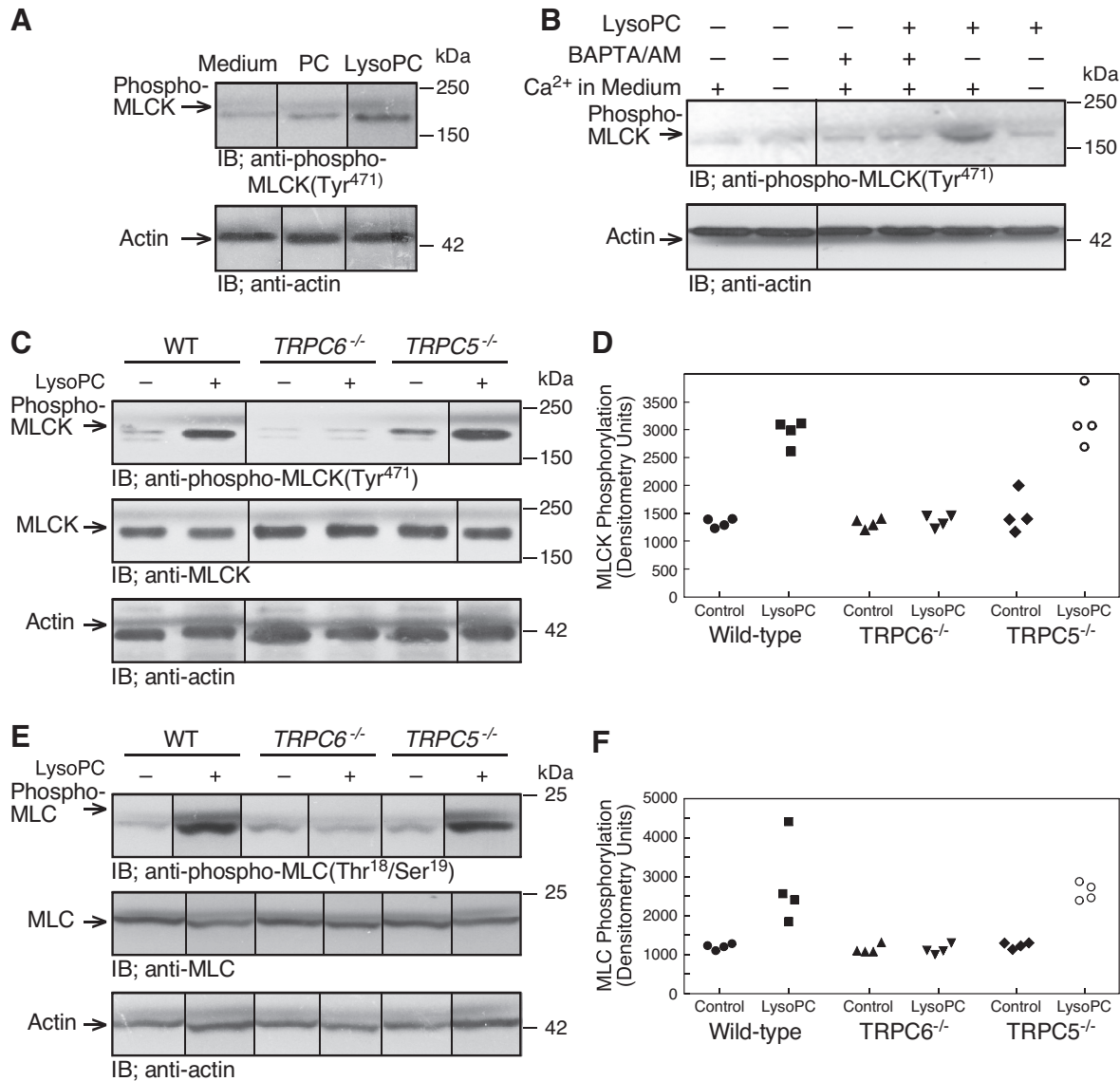


Fig. 6. LysoPC-induced MLCK phosphorylation is dependent on Ca^{2+} and TRPC6 channels. *A*: ECs were incubated with PC (12.5 μ M) or lysoPC (12.5 μ M) for 15 min. Phospho-MLCK(Tyr⁴⁷¹) was detected by immunoblot analysis, and actin served as loading control ($n = 3$). *B*: ECs were incubated in calcium-free or calcium-containing KR buffer for 10 min or ECs were incubated with BAPTA/AM (25 μ M) for 30 min before adding lysoPC. Phospho-MLCK(Tyr⁴⁷¹) was detected by immunoblot analysis, and actin served as loading control ($n = 4$). *C–F*: WT, TRPC5^{-/-}, or TRPC6^{-/-} MAECs were incubated with lysoPC (10 μ M) for 15 min. Phospho-MLCK(Tyr⁴⁷¹) and total MLCK (*C*) or phospho-MLC(Thr¹⁸/Ser¹⁹) and total MLC (*E*) were detected by immunoblot analysis. Actin served as loading control ($n = 4$). Densitometry measurements of phospho-MLCK(Tyr⁴⁷¹) (*D*) or phospho-MLC(Thr¹⁸/Ser¹⁹) (*F*) are represented in graphic form: ●, medium control with WT ECs; ■, lysoPC with WT ECs; ▲, medium control with TRPC6^{-/-} ECs; ▼, lysoPC with TRPC6^{-/-} ECs; ◆, medium control with TRPC5^{-/-} ECs; ○, lysoPC with TRPC5^{-/-} ECs. (In *A–C* and *E*, lines indicate lanes rearranged from the same gel. All bands are from the same gel.)

MLCK, the effect of MLCK downregulation using siRNA on EC migration was also assessed. Results using MLCK siRNA paralleled those with ML-9. Basal migration of control ECs and ECs transfected with NsiRNA or MLCK siRNA was similar. LysoPC inhibited 24-h migration of ECs transfected with NsiRNA by 64%, but in ECs transiently transfected with MLCK siRNA, lysoPC inhibited EC migration by only 43% ($n = 3$; $P < 0.001$, compared with ECs transfected with NsiRNA and incubated with lysoPC; Fig. 8*H*). This preservation of migration was similar to that seen in TRPC5^{-/-} MAECs or when TRPC5 was downregulated using siRNA (5, 33). Taken together, these data suggested that MLCK is essential in the lysoPC-induced TRPC5 activation and inhibition of EC migration.

DISCUSSION

TRPC5 is activated by a variety of agonists and through multiple mechanisms (50). Ca^{2+} -dependent regulation of Ca^{2+} -permeable cation channels is a ubiquitous mechanism of regulation. We have shown that lysoPC, the major lysophospholipid in oxidized LDL, activates TRPC6 with a resultant increase in $[Ca^{2+}]_i$ and this sets in motion a cascade of events resulting in TRPC5 externalization with a prolonged increase in $[Ca^{2+}]_i$ that disrupts the coordinated calcium transients required for normal EC migration. The studies presented here define some of the steps in this cascade.

An increase in $[Ca^{2+}]_i$ via TRPC6 channels can activate several pathways and kinases including ERK (18). Ca^{2+} and CaM can regulate the Ras/Raf/MEK/ERK pathway and posi-

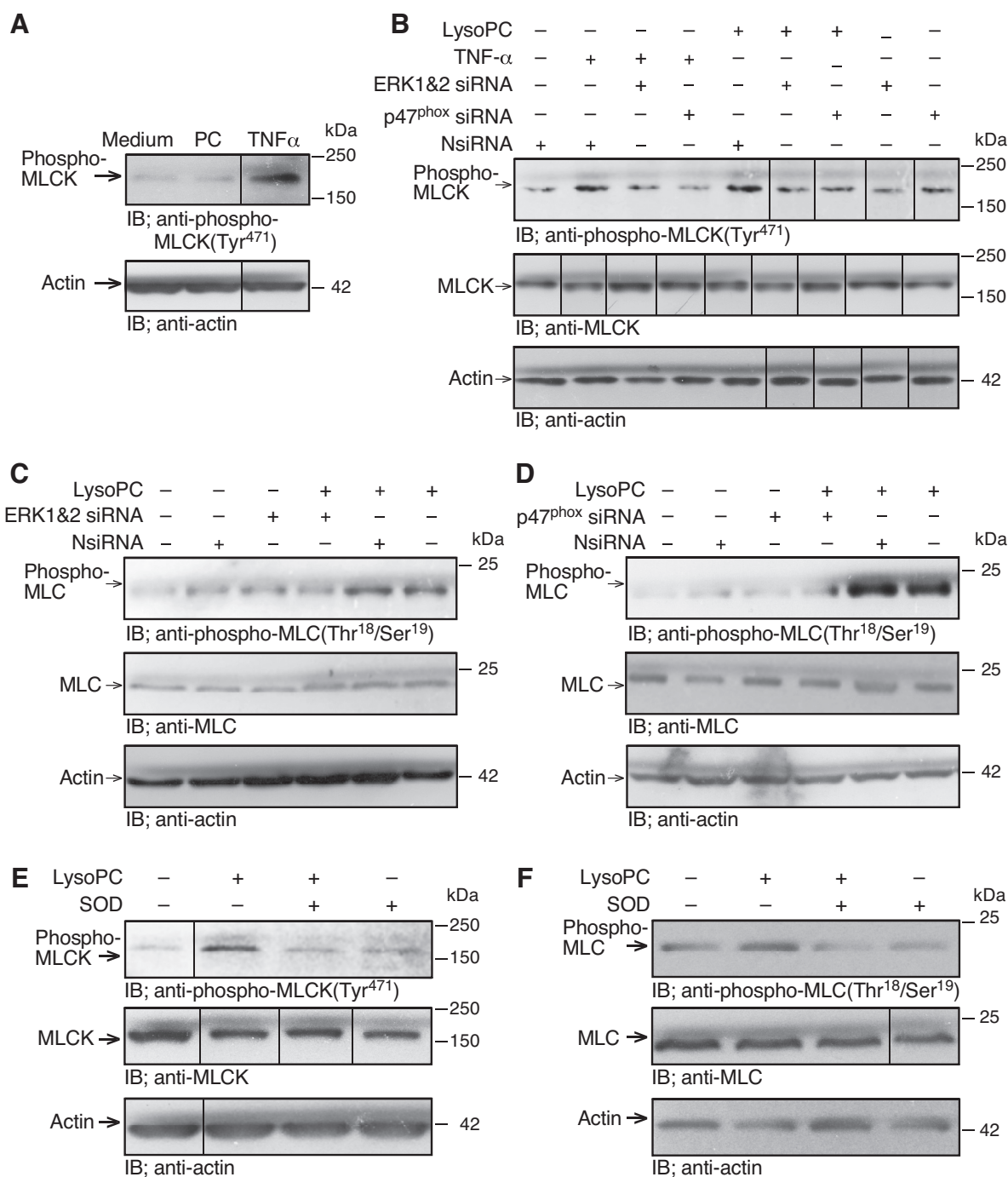


Fig. 7. LysoPC-induced MLCK and MLC phosphorylation are dependent on ERK and NADPH oxidase. *A*: ECs were incubated with PC (12.5 μ M) or TNF- α (500 U/ml) for 30 min. Phospho-MLCK(Tyr⁴⁷¹) was detected by immunoblot analysis, and actin served as loading control ($n = 3$). *B–D*: ECs were transiently transfected with NsiRNA (40 nM), ERK1 siRNA and ERK2 siRNA (20 nM of each), or p47^{phox} siRNA (20 nM) for 24 h before incubation with lysoPC or TNF- α . *B*: phospho-MLCK(Tyr⁴⁷¹) and total MLCK were detected by immunoblot analysis, and actin served as loading control ($n = 4$). *C* and *D*: phospho-MLC(Thr¹⁸/Ser¹⁹) and total MLC were detected by immunoblot analysis, and actin served as loading control ($n = 3$). *E* and *F*: ECs were incubated with superoxide dismutase (50 U/ml) for 30 min, then lysoPC (12.5 μ M) was added for 15 min. *E*: phospho-MLCK(Tyr⁴⁷¹) and total MLCK were detected by immunoblot analysis. Actin served as loading control ($n = 3$). *F*: phospho-MLC(Thr¹⁸/Ser¹⁹) and total MLC were detected by immunoblot analysis. Actin served as loading control ($n = 3$). (In *A* and *B*, *E* and *F*, lines indicate lanes rearranged from the same gel. All bands are from the same gel.)

tively modulate ERK1/2 activation (1, 24). We show that lysoPC-induced ERK activation is regulated by influx of Ca²⁺ specifically through TRPC6 channels. ERK and TRPC5 are not activated when ECs from *TRPC6*^{-/-} mice are exposed to

lysoPC. These findings are consistent with the work of other investigators showing increased ERK1/2 phosphorylation in 293T cells and podocytes that overexpress TRPC6 proteins with gain-of-function mutations (10).

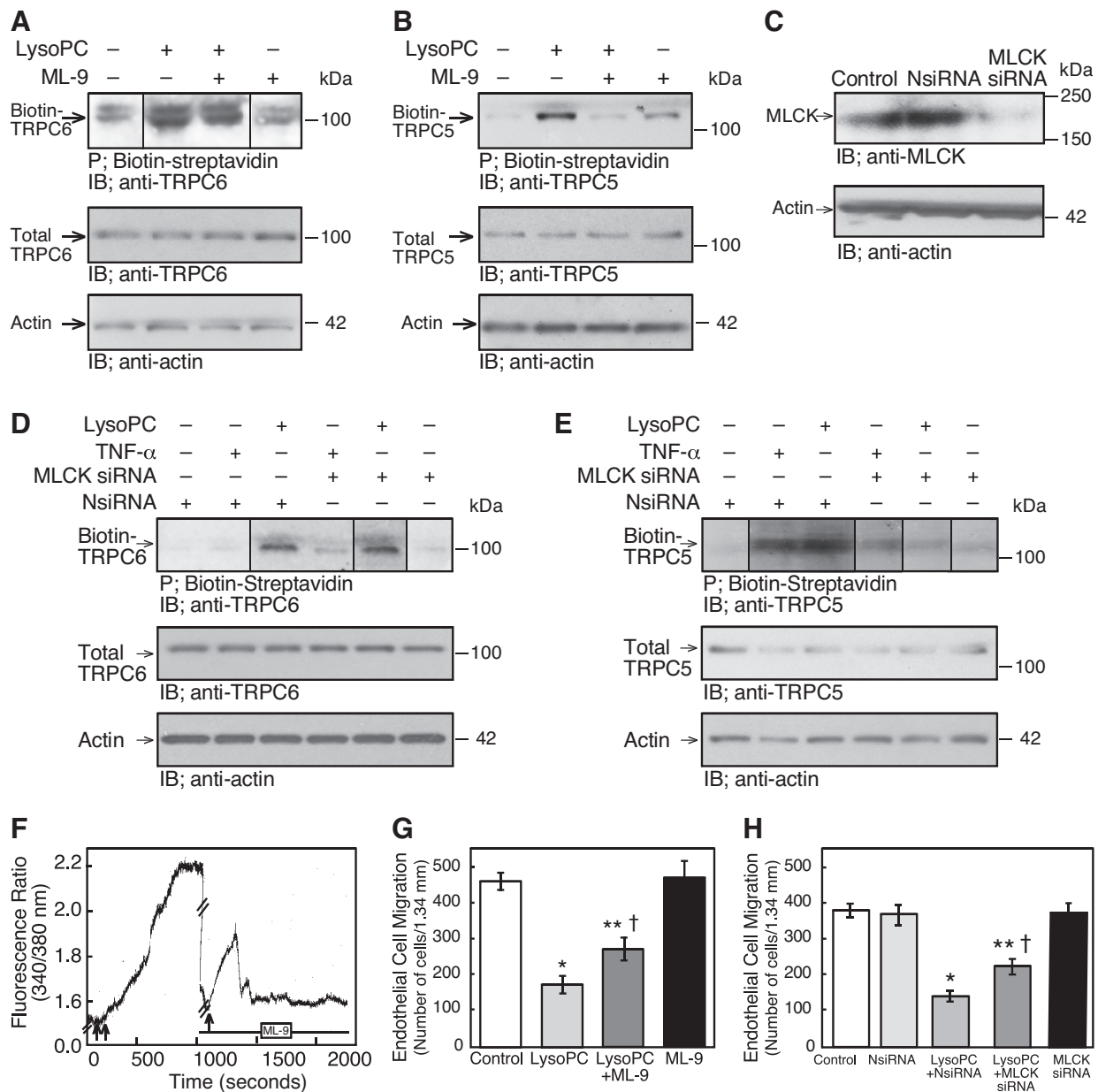


Fig. 8. MLCK inhibition blocks lysoPC-induced TRPC5 externalization and increased $[Ca^{2+}]_i$, and partially preserves EC migration in lysoPC. *A* and *B*: ECs were pretreated with ML-9 (10 μ M) for 30 min before incubation with lysoPC (12.5 μ M). Externalization of TRPC6 or TRPC5 was detected by biotinylation assay and total TRPC6 or TRPC5 by immunoblot analysis. Actin served as loading control ($n = 4$). *C*: ECs were transiently transfected with NsiRNA or MLCK siRNA (30 nM) for 24 h. The siRNA was removed, and 24 h later the cells were lysed. MLCK was detected by immunoblot analysis. Actin served as loading control ($n = 4$). *D* and *E*: ECs were transiently transfected with NsiRNA or MLCK siRNA for 24 h before incubation with lysoPC or TNF- α . Externalization of TRPC6 or TRPC5 was detected by biotinylation assay and total TRPC6 or TRPC5 by immunoblot analysis. Actin served as loading control ($n = 4$). *F*: ECs were made quiescent for 18 h, then loaded with fura 2-AM. After the baseline $[Ca^{2+}]_i$ was established, lysoPC (12.5 μ M) was added to the cells (double arrow) and relative change of $[Ca^{2+}]_i$ was measured. After $[Ca^{2+}]_i$ plateaued, lysoPC was removed by washing with KR buffer. After readings returned to baseline (in 7–10 min), ML-9 (10 μ M) was added and recordings made for 60 s. After 20 min, lysoPC (12.5 μ M, arrow) was added, and the relative change of $[Ca^{2+}]_i$ was measured. The tracing is representative of four separate experiments (///denotes the wash period and ML-9 incubation period during which times no data were collected). *G*: ECs were pretreated with ML-9 (10 μ M) for 30 min, then the ML-9 was removed, and the migration assay initiated. Migration was assessed after 24 h in the presence or absence of lysoPC. Results are represented as means \pm SD ($n = 5$; * $P < 0.001$, compared with control; ** $P < 0.001$, compared with ML-9 control; † $P < 0.001$, compared with lysoPC). *H*: ECs were transiently transfected for 24 h with NsiRNA or MLCK siRNA (30 nM) and then made quiescent for 12 h. The migration assay initiated, and migration in the presence or absence of lysoPC was assessed after 24 h. Results were represented as means \pm SD ($n = 5$; * $P < 0.001$, compared with control; ** $P < 0.001$, compared with MLCK siRNA; † $P < 0.001$, compared with lysoPC with NsiRNA). (In *A*, *D* and *E*, lines indicate lanes rearranged from the same gel. All bands are from the same gel.)

We have shown that lysoPC increases EC production of ROS through activation of NADPH oxidase and this contributes to the inhibition of EC migration (43). Activation of NADPH oxidase in ECs involves phosphorylation and translocation of cytosolic components, including p47^{phox} and p67^{phox}, to the membrane where they assemble with Nox2 or Nox4 and p22^{phox} (29). NADPH oxidase activation by specific agonists requires an increase in intracellular calcium (11), and lysophosphatidylcholines prime NADPH oxidase and stimulate multiple neutrophil functions through changes in cytosolic Ca²⁺ (38). We provide evidence that the lysoPC-induced activation of NADPH oxidase in ECs depends on Ca²⁺ influx through TRPC6-containing channels because lysoPC does not activate NADPH oxidase in *TRPC6*^{-/-} MAECs. TRPC6 can be forming a complex with other TRPC proteins to permit Ca²⁺ influx to activate NADPH oxidase as has been shown in granulocytes (4). Furthermore, the resultant Ca²⁺ influx can activate NADPH oxidase directly or indirectly through activation of a kinase.

ERK can activate NADPH oxidase (13, 35). When ECs are exposed to transient hypoxia, ERK2 is activated, and this in turn activates NADPH oxidase causing an increase in ROS production (35). Our finding that downregulation of ERK blocks lysoPC-induced NADPH activation also indicates that ERK activation is upstream from NADPH oxidase activation. We find that downregulation of p47^{phox} or p22^{phox} does not alter lysoPC-induced ERK phosphorylation. Similarly, angiotensin II-induced ROS formation is blocked by p22^{phox} antisense, but this has no effect on angiotensin II-induced ERK activation (44). Our results suggest that Ca²⁺ influx through TRPC6 channels activates ERK that in turn activates NADPH oxidase.

The importance of NADPH oxidase activation lies, in part, in the ability of superoxide to function as an intracellular signaling molecule. Although ROS can activate MLCK (51), and MLCK can activate TRPC5 channels (21, 37), there is conflicting evidence in the literature on the ability of ROS to activate TRPC5. One report indicates that TRPC5 can be activated by ROS (27). Another study suggests that ROS do not activate TRPC5, but only one inhibitor was used and its effectiveness was not assessed (16). Our findings support a major role for ROS in the lysoPC-induced externalization of TRPC5.

ERK can phosphorylate MLCK and thereby increase its activity (3, 22), and MLCK is essential for TRPC5 channel externalization (21, 37). MLCK phosphorylates MLC, and MLC phosphorylation/dephosphorylation regulates the insertion of TRPC5 containing vesicles into the plasma membrane (21, 37). Intracellular Ca²⁺-calmodulin may constitutively activate MLCK maintaining a basal distribution of TRPC5 channels at plasma membrane (37). Our data suggest that lysoPC increases MLCK activity through ERK activation and subsequent MLCK phosphorylation.

Inhibition of MLCK impairs TRPC5 translocation to the plasma membrane and TRPC5 activity (21, 37). We find that ML-9 can inhibit the lysoPC-induced externalization of TRPC5 channels in ECs, complementing the findings in HEK293 cells overexpressing TRPC5 where ML-9 decreased the cell surface localization of TRPC5 channels (37). In HEK293 cells overexpressing TRPC6, ML-9 has been reported to reduce cationic currents elicited by carbachol by an MLCK-independent mechanism (36). Our data do not show a reduction in basal [Ca²⁺]_i in ECs pretreated with ML-9 nor do the results reported by other investigators studying ECs (39, 45, 46). The ability of ML-9 to reduce [Ca²⁺]_i may be dependent on concentration, cell type, and native expression of calcium channels.

The pattern of lysoPC-induced increase in [Ca²⁺]_i seen in ECs pretreated with ML-9 is very similar to that seen after blocking of TRPC5 activity with siRNA (5). In both cases, lysoPC induces only an initial transient increase in [Ca²⁺]_i that we attribute to TRPC6 channels, but the prolonged elevation of [Ca²⁺]_i is absent. ML-9 inhibits the lysoPC-induced externalization of TRPC5 but not TRPC6, supporting the role of MLCK in TRPC5, but not TRPC6, insertion into the cell membrane. LysoPC-induced phosphorylation of MLCK is blocked in ECs from *TRPC6*^{-/-} mice but not in those from *TRPC5*^{-/-} mice, suggesting that Ca²⁺ influx through TRPC6 is critical for lysoPC-induced MLCK activation.

Our findings should not be taken to suggest that there is a single linear pathway between TRPC6 channel opening and TRPC5 channel externalization. Many alternative pathways, redundancies, and regulatory influences exist. ROS production by NADPH oxidase can activate ERK (12, 49). MLCK can facilitate recruitment of cytoskeletal proteins and NADPH oxidase to the plasma membrane with subsequent activation of

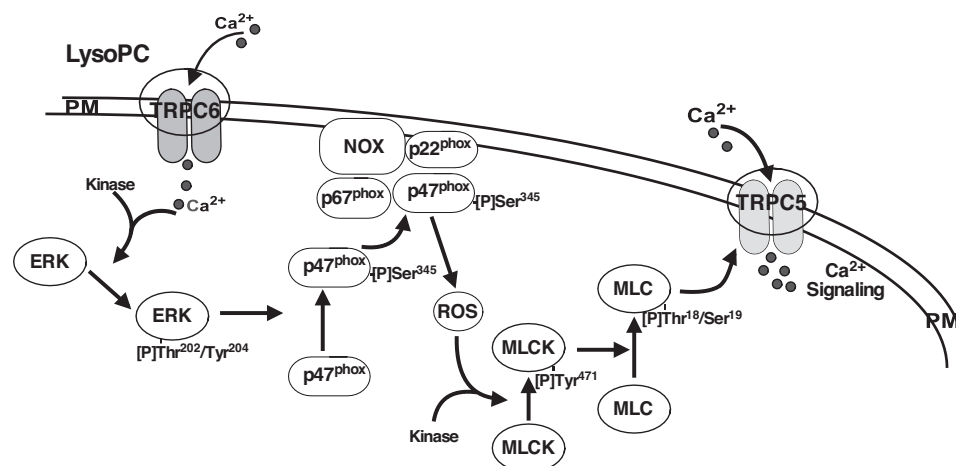


Fig. 9. Proposed signaling pathway between TRPC6 activation and TRPC5 externalization. Ca²⁺ entry through TRPC6 activates a kinase, leading to ERK phosphorylation. ERK then phosphorylates p47^{phox}, leading to NADPH oxidase activation and production of ROS, which contribute to the phosphorylation of MLCK that is required for externalization of TRPC5. The components of this process may be localized in a “signalsome” of the plasma membrane (PM).

NADPH oxidase and increased ROS production (42). Pathologic feed-forward activities could occur with cycles of increasing ROS production and ERK and MLCK activation, culminating in elevated TRPC5 externalization.

The in vivo relevance of these studies is seen in comparing endothelial healing of arterial injuries in mice on a high-cholesterol diet compared with those on a chow diet (34). The high-cholesterol diet increases plasma lysoPC levels and inhibits endothelial healing in WT mice. Although a high-cholesterol diet causes the same increase of plasma cholesterol and a greater increase of plasma lysoPC in *TRPC6*^{-/-} and *TRPC5*^{-/-} mice, endothelial healing is not impaired in *TRPC6*^{-/-} mice and the impairment is significantly reduced in *TRPC5*^{-/-} mice (33). These findings suggest that the presence or absence of TRPC6 and TRPC5 channels has a significant influence on the cellular response to environmental factors.

Based on our results, we propose that lysoPC induces CaM-dependent TRPC6 activation allowing an initial transient rise in [Ca²⁺]_i. This triggers ERK activation, which leads to the activation of NADPH oxidase. The increase in ROS activates MLCK, which in turn facilitates TRPC5 channel insertion into the plasma membrane and causes a prolonged influx of Ca²⁺ that disrupts the finely coordinated local signals that regulate calcium transients required for normal EC migration (see proposed model in Fig. 9). Although studies were conducted using animal ECs, TRPC6 and TRPC5 are expressed in human ECs, and downregulation of TRPC6 blocks lysoPC-induced p47^{phox} phosphorylation (Fig. 4D), suggesting that the pathway outlined in this paper is functional in human ECs. Knowledge of the signaling pathway provides opportunities for its regulation and the restoration of endothelial healing under conditions of elevated lipid oxidation products or oxidative stress leading to TRPC6 or TRPC5 activation.

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DISCLOSURES

No conflicts of interest, financial or otherwise, are declared by the authors.

AUTHOR CONTRIBUTIONS

P.C., L.B., and L.M.G. conceived and designed research; P.C. and M.A.R. performed experiments; P.C., M.A.R., and L.M.G. analyzed data; P.C., M.A.R., and L.M.G. interpreted results of experiments; P.C., M.A.R., and L.M.G. prepared figures; P.C. and L.M.G. drafted manuscript; P.C., M.A.R., L.B., and L.M.G. edited and revised manuscript; P.C., M.A.R., L.B., and L.M.G. approved final version of manuscript.

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