Caveolin-1 Assembles Type 1 Inositol 1,4,5-Trisphosphate Receptors and Canonical Transient Receptor Potential 3 Channels into a Functional Signaling Complex in Arterial Smooth Muscle Cells

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Physical coupling of sarcoplasmic reticulum (SR) type 1 inositol 1,4,5-trisphosphate receptors (IP₃R1) to plasma membrane canonical transient receptor potential 3 (TRPC3) channels activates a cation current (I_cat) in arterial smooth muscle cells that induces vasoconstriction. However, structural components that enable IP₃R1 and TRPC3 channels to communicate locally are unclear. Caveolae are plasma membrane microdomains that can compartmentalize proteins. Here, we tested the hypothesis that caveolae and specifically caveolin-1 (cav-1), a caveolae scaffolding protein, facilitate functional IP₃R1 to TRPC3 coupling in smooth muscle cells of resistance-size cerebral arteries. Methyl-β-cyclodextrin (MβCD), which disassembles caveolae, reduced IP₃R1 activation in smooth muscle cells and vasoconstriction in pressurized arteries. Cholesterol replenishment reversed these effects. Cav-1 knockdown using shRNA attenuated IP₃R1-induced vasoconstriction, but did not alter TRPC3 and IP₃R1 expression. A synthetic peptide corresponding to the cav-1 scaffolding domain (CSD) sequence (amino acids 82–101) also attenuated IP₃R1-induced I_cat activation and vasoconstriction. A cav-1 antibody co-immunoprecipitated cav-1, TRPC3, and IP₃R1 from cerebral artery lysate. ImmunoFRET indicated that cav-1, TRPC3 channels and IP₃R1 are spatially co-localized in arterial smooth muscle cells. IP₃R1 and TRPC3 channel spatial localization was disrupted by MβCD and a CSD peptide. Cholesterol replenishment re-established IP₃R1 and TRPC3 channel close spatial proximity. Taken together, these data indicate that in arterial smooth muscle cells, cav-1 co-localizes SR IP₃R1 and plasma membrane TRPC3 channels in close spatial proximity thereby enabling IP₃R1-induced physical coupling of these proteins, leading to I_cat generation and vasoconstriction.

Several G-protein-coupled receptor agonists activate phospholipase C leading to generation of the second messenger, inositol 1,4,5-trisphosphate (IP₃)² (1). IP₃ binds to endo/sarcoplasmic reticulum-localized IP₃ receptors leading to Ca²⁺ release from the endo/sarcoplasmic reticulum (2). In arterial smooth muscle cells, the resultant increase in intracellular Ca²⁺ concentration ([Ca²⁺]) causes vasoconstriction. IP₃ also activates a non-selective cation current (I_cat), leading to vasoconstriction (3–7). IP₃-induced I_cat activation occurs independently of sarcoplasmic reticulum (SR) Ca²⁺ release and due to a physical interaction between type 1 IP₃ receptors (IP₃R1) and canonical transient receptor potential (TRPC) 3 channels (4–6). IP₃R1 selectively couples to TRPC3 channels due to close spatial proximity of these proteins (4). A wide variety of signal transduction pathways require interacting molecules to form functional macromolecular complexes. These complexes often contain one or more scaffolding proteins that traffic, localize, or compartmentalize signaling molecules (8). However, whether scaffolding proteins are required to assemble IP₃R1 and TRPC3 channels into a functional macromolecular complex in arterial smooth muscle cells is unclear.

Caveolae are plasma membrane microdomains that act as compartments for several signal transduction molecules, including receptors, ion channels, and scaffolding proteins (9). Caveolin integrity is sustained by cholesterol and caveolin proteins, of which there are three different isoforms (cav-1–3). Caveolin isoforms can exhibit tissue-specific expression (9). Cav-1 is required for caveoleae formation in arterial smooth muscle cells, embryonic fibroblasts, lung endothelial, and epithelial cells (10–13). Cav-1 ablation and caveolae disruption attenuate both pressure- and agonist-induced vasoconstriction, indicating that caveolae play an important role in the regulation of vascular reactivity (14, 16). In arterial smooth muscle cells, peripheral SR membrane can be located nearby plasma membrane caveolae (16). This close spatial proximity may permit or enhance interactions between SR- and caveole-localized plasma membrane proteins. Whether IP₃R1, TRPC3 channels, and caveolin proteins physically and functionally interact in arterial smooth muscle cells is unclear.
Caveolin-1 Assembles IP₃R1-TRPC3 Signaling Complex

In the present study, we examined the significance of caveolae in mediating IP₃-induced I_{Ca,ATP} activation in cerebral artery smooth muscle cells and vasoconstriction in pressurized cerebral arteries. We show that caveolae disassembly using cycloheximide, cave-1 knockdown using RNAi, or a cave-1 scaffolding domain (CSD) peptide that competes with endogenous cave-1 for interacting protein binding, inhibit IP₃-induced I_{Ca,ATP} activation in smooth muscle cells and vasoconstriction. Data indicate that IP₃R1 and TRPC3 physically interact with cave-1, and that cave-1 and caveolae maintain plasma membrane TRPC3 and SR IP₃R1 in a functional signaling complex in cerebral artery smooth muscle cells, thereby permitting IP₃-induced vasoconstriction.

EXPERIMENTAL PROCEDURES

Tissue Preparation and Smooth Muscle Cell Isolation—Animal protocols used were reviewed and approved by the Animal Care and Use Committee at the University of Tennessee Health Science Center. Male Sprague-Dawley rats (6–8 weeks) were euthanized by intraperitoneal injection of sodium pentobarbital (150 mg/kg). The brain was removed and resistance-size cerebral arteries (posterior cerebral, cerebral, and middle cerebral, <200 μm) were harvested and maintained in ice-cold (4 °C), oxygenated (21% O₂-5% CO₂-74% N₂) physiological saline solution (PSS) containing (in mM): 119 NaCl, 4.7 KCl, 24 NaHCO₃, 1.2 KH₂PO₄, 1.6 CaCl₂, 1.2 MgSO₄, and 11 glucose (pH 7.4). Individual smooth muscle cells were dissociated from cerebral arteries using a HEPES-buffered solution containing (in mM): 55 NaCl, 80 sodium glutamate, 5.6 KCl, 2 MgCl₂, 10 HEPES, and 10 glucose (pH 7.3), as previously described (18). Briefly, cerebral arteries were placed into isolation solution containing 0.7 mg/ml papain, 1 mg/ml dithioerythreitol, and 1 mg/ml bovine serum albumin (BSA) for 12 min at 37 °C. Arteries were then immediately transferred to isolation solution containing 1 mg/ml collagenase F and H (2:1), 10 μM CaCl₂, and 1 mg/ml BSA for 10 min at 37 °C. Arteries were subsequently washed in isolation solution and triturated using a fire-polished glass pipette to yield single smooth muscle cells. Cells were maintained at 4 °C, and used for experimentation within 8 h.

Patch-Clamp Electrophysiology—Membrane currents and cell capacitance were measured in isolated smooth muscle cells using the patch-clamp technique (Axopatch 200B, Clampex 8.2). I_{Ca,ATP} was measured using the conventional whole cell patch-clamp configuration, as we have done previously (5, 6). Briefly, whole cell currents were measured by applying 940-ms voltage ramps between −120 and +20 mV from a holding potential of −40 mV. Bath solution for conventional whole-cell experiments contained (in mM): 140 NaCl, 1.8 CaCl₂, 1.2 MgCl₂, 10 HEPES, and 10 glucose (pH 7.4). The pipette solution contained (in mM): 140 CsCl, 10 HEPES, 10 glucose, 5 Mg-ATP, and 5 EGTA (with pH adjusted to 7.2 with CsOH), and 100 mM free Ca²⁺.

Pressurized Artery Diameter Measurement—Arterial diameter was measured using pressurized artery myography, as previously described (19). Briefly, an arterial segment 1–2 mm in length was cannulated at each end in a pressurized artery chamber (Living Systems Instrumentation; Burlington, VT). The chamber was continuously perfused with PSS equilibrated with a mixture of 21% O₂, 5% CO₂, 74% N₂, and maintained at 37 °C. Arterial endothelium was denuded by introducing air bubbles into the artery lumen for 1 min followed by a wash with PSS. Intravascular pressure was changed using an attached reservoir and monitored using a pressure transducer. Pressurized arteries were observed with a charge-coupled device camera attached to an inverted microscope (Nikon TE 200). Arterial diameter was measured at 1 Hz using the automatic edge-detection function of IonWizard software (Ionoptix; Milton, MA). Pharmacological agents were applied via chamber perfusion after arteries had developed steady-state spontaneous myogenic tone. Myogenic tone (%) was calculated as (1 − active diameter/passive diameter) × 100.

Cav-1 Knockdown—Using pRNA-U6.1/Neo as a template, silencing vectors were constructed to express short hairpin (sh) RNA (GenScript Corp, Piscataway, NJ) as previously described (5). Expressed DNA sequences were as follows: cav-1shV1: AAGCATCTCAACGACGACGTG, cav-1shV2: AAGACTGCAGATTGCGTTTG, cav-1scrm: GCACAGC-AAGGATCATGCGTGA. The two silencing vectors (cav-1shV1 + cav-1shV2; cav-1shV) or a scrambled control (cav-1scrm) shRNA were inserted into cerebral artery segments using a reversible permeabilization procedure, as previously described (20). Arteries were placed into serum-free Dulbecco’s modified Eagle’s medium (DMEM) supplemented with 1% penicillin/streptomycin and incubated at 37 °C (95% O₂, 5% CO₂) for 4 days.

Co-immunoprecipitation and Western Immunoblotting—Arterial lysate (yielding ~1.5 mg total protein) was harvested from cerebral arteries using ice-cold non-denaturing lysis buffer (Thermo Scientific, Rockford, IL). Because of the small size (<200 μm diameter) of the cerebral arteries used in this study, arteries pooled from ~15 rats were required for each co-IP experiment. Co-immunoprecipitation (co-IP) was done using the Thermo Scientific Pierce co-IP kit following the manufacturer’s protocol. Briefly, the cav-1 antibody (Abcam) was first immobilized for 2 h using AminoLink Plus coupling resin. The resin was then washed and incubated with arterial lysate overnight. After incubation, the resin was again washed and protein eluted using elution buffer. A negative control that was provided with the IP kit to assess nonspecific binding received the same treatment as the co-IP samples, including the cav-1 antibody. In this control, the coupling resin is not amine-reactive preventing covalent immobilization of the primary antibody onto the resin. Samples were analyzed by Western blotting using rabbit polyclonal anti-cav-1 (Abcam Inc., Cambridge, MA), mouse monoclonal anti-IP₃R1 (UC Davis/NINDS/NIMH NeuroMab Facility), rabbit polyclonal anti-TRPC3 (Alomone Labs Ltd. Jerusalem, Israel), and horseradish peroxidase-conjugated secondary antibodies.

Western blotting was performed as previously described (4–6). Briefly, rat cerebral artery lysate protein concentrations were determined spectrophotometrically. Proteins were separated by 7.5% gradient SDS-polyacrylamide gel electrophoresis and transferred onto nitrocellulose membranes using a Mini Trans Blot Cell (Bio-Rad). Membranes were then incu-
bated with respective antibodies and developed using enhanced chemiluminescence (Thermo Scientific).

**Chemical Loading**—Caveolin-1 scaffolding domain peptide (CSD) was inserted into intact cerebral arteries using the chemical loading technique modified from a previously described method (14, 21). Briefly, cerebral arteries were first placed into a Ca\(^{2+}\)-free (EGTA, 10 mM), high K\(^+\) (120 mM) solution supplemented with ATP (5 mM) at 4 °C for ~30 min. Arteries were then incubated in the presence of CSD in Mg\(^{2+}\) (2 mM)- and ATP (5 mM)-containing Ca\(^{2+}\)-free solution at 4 °C for ~2 h. Arteries were further placed in 10 mM Mg\(^{2+}\) and ATP (5 mM)-containing Ca\(^{2+}\)-free solution at 4 °C for ~30 min, after which the solution was changed to one containing Na\(^+\) (140 mM) and K\(^+\) (5 mM) and maintained at room temperature for 30 min. Ca\(^{2+}\) was then added incrementally to the solution to reach a concentration of 1.8 mM. Arteries were then placed into DMEM-F12 supplemented with 1% penicillin and streptomycin overnight (37 °C, 5% CO\(_2\)) prior to use.

**Immunofluorescence Resonance Energy Transfer (ImmunofRET) and Confocal Imaging**—ImmunofRET was performed using a modified version of our previously described method (4). Briefly, paraformaldehyde-fixed cerebral artery smooth muscle cells were permeabilized with 0.1% Triton X-100 for 1 min at room temperature. Following a 1 h incubation in PBS containing 5% bovine serum albumin (BSA), smooth muscle cells were treated overnight at 4 °C with mouse monoclonal anti-IP3R1 (UC Davis/NINDS/NIMH NeuroMab Facility), sheep polyclonal anti-TRPC3 (Abcam), rabbit polyclonal anti-cav-1 (Abcam), or anti-IP3R1 plus anti-TRPC3 or anti-TRPC3 plus anti-cav-1 at a dilution of 1:100 each in PBS containing 5% BSA. After a wash and block with PBS containing 5% BSA, cells were incubated for 1 h at 37 °C with secondary antibodies: Alexa 546-conjugated donkey anti-mouse for IP3R1 (1:100 dilution; Invitrogen, Carlsbad, CA) and Alexa 488-conjugated donkey anti-sheep for TRPC3 (1:100 dilution; Invitrogen). For FRET measurements using TRPC3 and cav-1 primary antibodies, Alexa 546-conjugated donkey anti-sheep and Alexa 488-conjugated goat anti-rabbit secondary antibodies were used, respectively. Following wash and mount, fluorescence images were acquired using a Zeiss LSM Pascal laser-scanning confocal microscope. Alexa 488 and Alexa 546 were excited at 488 and 543 nm and emission collected at 505–530 nm. Negative controls prepared without primary antibodies and developed using enhanced chemiluminescence did not exhibit fluorescence. Images were background-subtracted and normalized FRET (N-FRET) was calculated on a pixel-by-pixel basis for the entire image and in regions of interest (within the boundaries of the cell) using the Xia method (22) and Zeiss LSM FRET Macro tool version 2.5.

Fluorescein isothiocyanate (FITC)-labeled antennapedia peptide was inserted into smooth muscle cells of intact cerebral arteries using chemical loading procedure. FITC fluorescence was background-subtracted. Control arteries underwent the same chemical loading and culture protocol, but were not exposed to FITC-labeled antennapedia. FITC fluorescence was imaged using a Zeiss LSM5 Pascal laser-scanning confocal microscope. FITC was excited at 488 nm and emission collected at 505–530 nm.

**Chemicals**—Unless otherwise stated, all reagents were purchased from Sigma. Papain was purchased from Worthington Biochemical (Lakewood, NJ) and 2,3,6-tri-O-butyryl-myoinositol 1,4,5-trisphosphate-hexakis (propionoxymethyl) ester (Bt-IP3) was purchased from SiChem (Bremen, Germany). Antennapedia and FITC-conjugated antennapedia peptides were purchased from Anaspec (Fremont, CA). Caveolin-1 scaffolding domain peptide was purchased from Enzo Life Sciences (Farmingdale, NY).

**Statistical Analysis**—Data are expressed as means ± S.E. of the mean. Statistical significance was calculated by using Student’s t-tests for paired or unpaired data and analysis of variance followed by Student-Newman-Keuls post-hoc test for multiple data sets. p < 0.05 was considered significant.

**RESULTS**

Caveolae Are Required for IP\(_3\)-induced I\(_{\text{cat}}\) Activation in Arterial Smooth Muscle Cells and Vasoconstriction—To examine the significance of caveolae for IP\(_3\)-induced I\(_{\text{cat}}\) activation, freshly-isolated cerebral artery smooth muscle cells were exposed to methyl β-cyclodextrin (MβCD; 30 min), a cholesterol-depleting agent that disassembles caveolae (15, 23). MβCD treatment reduced mean smooth muscle cell capacitance (control versus MβCD-treated (pF): 12.1 ± 1.0, n = 6 versus 7.4 ± 1.2, n = 7; p < 0.05), consistent with a decrease in cell surface area. MβCD treatment also reduced IP\(_3\)-induced I\(_{\text{cat}}\) density in arterial smooth muscle cells by ~89% (Fig. 1, A and B). To study the importance of cholesterol in
mediating this effect, cholesterol-loaded MβCD (Chol/MβCD; 1 h) was used to replenish cholesterol in MβCD-treated arterial smooth muscle cells (15, 23, 24). Cholesterol replenishment reversed the MβCD-induced decrease in cell capacitance (to 10.7 ± 0.9 pF, n = 6, p < 0.05 when compared with control). Chol/MβCD also reversed the inhibitory effect of MβCD on IP3-induced ICaL activation (Fig. 1, A and B).

The functional significance of caveolae for IP3-induced vasconstriction was investigated by measuring diameter regulation of pressurized (60 mmHg), endothelium-denuded cerebral arteries. Control arteries developed 25% mean myogenic tone (Fig. 2, A and B). MβCD treatment (2 h) reduced myogenic tone to 7%, or by 72% (Fig. 2, A and B). MβCD treatment also reduced vasoconstriction induced by Bt-IP3, a membrane-permeant IP3 analog, by 79% (Fig. 2, C and D). Chol/MβCD treatment (~2 h) restored both myogenic tone and IP3-induced vasoconstriction in arteries treated with MβCD (Fig. 2, A–D). These data suggest that intact caveolae are necessary for IP3-induced ICaL activation in arterial smooth muscle cells and myogenic tone and IP3-induced vasoconstriction in pressurized arteries.

Cav-1 Expression Is Necessary for IP3-induced Vasoconstriction—Cav-1 is essential for caveolae formation in cerebral artery smooth muscle cells (13). Therefore, to study the importance of cav-1 in IP3-induced vasoconstriction, vectors encoding shRNA targeting cav-1 (cav-1shV) were inserted into cerebral arteries to suppress cav-1 expression. Western blotting indicated that cav-1shV reduces mean arterial cav-1 protein by 41% when compared with arteries treated with scrambled shRNA (scrm; Fig. 3, A and B). In contrast, cav-1shV did not alter TRPC3 or IP3R1 expression (Fig. 3, A and B). Cav-1 knockdown reduced myogenic tone in endothelium-denuded pressurized (60 mmHg) arteries by 50% (Fig. 4, A and B). Cav-1 knockdown also reduced Bt-IP3-induced vasoconstriction by ~61% (Fig. 4, C and D). These data indicate that cav-1 expression is required for myogenic tone and IP3-induced vasoconstriction in cerebral arteries.

CSD Peptide Attenuates IP3-induced ICaL Activation and Vasoconstriction—The cav-1 N terminus contains a CSD that interacts with and regulates a wide variety of signaling proteins containing a CSD-interacting motif (25, 26). CSD-interacting motifs are characterized as X4X4X5XX, where X is a tryptophan, phenylalanine, or...
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tyrosine, and X is any amino acid (26). As shown in Fig. 5A, the N and C termini of IP$_3$R1 and the C terminus of TRPC3 channels contain CSD-binding motifs, suggesting that IP$_3$R1 and TRPC3 channels may both interact with cav-1.

To study the physiological function of the CSD in IP$_3$-induced I$_{Ca}^\text{L}$ activation and vasoconstriction, we used a synthetic peptide corresponding to the cav-1 CSD sequence (amino acids 82–101, DGIWKA$\text{SFTFTVTKYWFYR}$). This CSD peptide competes with endogenous cav-1 for interacting protein binding. CSD peptide was conjugated to an antenna-pedia (ANT) internalization sequence (ANT-CSD, amino acids 43–58, RQIKIWFQNRMKWKK) to improve membrane permeability. An ANT peptide was used as an experimental control. To examine membrane permeability of the peptide, a FITC-labeled ANT peptide (ANT-FITC) was used. Confocal imaging of arteries exposed to ANT-FITC indicated successful introduction into smooth muscle cells (Fig. 5B).

ANT-CSD peptide did not alter smooth muscle cell capacitance (ANT versus ANT-CSD, 12 ± 2 versus 11 ± 1 pF, n = 5 and 6, respectively, p > 0.05). However, ANT-CSD peptide reduced IP$_3$-induced I$_{Ca}^\text{L}$ density by ~48% when compared with control (ANT; Fig. 5, C and D). ANT-CSD peptide did not alter myogenic tone in endothelium-denuded pressurized (60 mmHg) cerebral arteries (ANT versus ANT-CSD, ~22 ± 6 versus 20 ± 3%, n = 6 and 5, respectively, p > 0.05). In contrast, ANT-CSD peptide attenuated IP$_3$-induced vasoconstriction by ~61% in pressurized (60 mmHg) cerebral arteries (Fig. 6, A and B). These data indicate that the CSD regulates IP$_3$-induced I$_{Ca}^\text{L}$ activation in arterial smooth muscle cells and IP$_3$-induced vasoconstriction in pressurized arteries.

Cerebral Artery cav-1 Immunoprecipitates with Both TRPC3 and IP$_3$R1—To test the hypothesis that cav-1 exists in a macromolecular protein complex with IP$_3$R1 and TRPC3 channels, co-IP was performed. A cav-1 antibody co-immunoprecipitated cav-1, IP$_3$R1, and TRPC3 from cerebral artery lysate (Fig. 7). These data indicate that cav-1, IP$_3$R1, and TRPC3 channels are located within the same macromolecular complex in cerebral arteries.

Cav-1 and TRPC3 Are Spatially Localized in Arterial Smooth Muscle Cells—Spatial localization of TRPC3 channels to cav-1 in arterial smooth muscle cells was studied using immunoFRET. Fig. 8A indicates that cav-1 and TRPC3 are co-localized in the arterial smooth muscle cell plasma membrane. Alexa 488- and Alexa 546-conjugated secondary antibodies bound to cav-1 and TRPC3 channel primary antibodies, respectively, generated mean N-FRET of 23.0 ± 1.2% (n = 16). These data indicate that cav-1 and TRPC3 are spatially localized in the cerebral artery smooth muscle cell plasma membrane.

IP$_3$R1 and TRPC3 Are in Close Spatial Proximity and Separated by Caveolae Disruption or CSD Peptide in Arterial Smooth Muscle Cells—ImmunoFRET was also used to measure the spatial proximity of IP$_3$R1 and TRPC3 channels...
in arterial smooth muscle cells. Alexa 488- and Alexa 546-labeled secondary antibodies bound to TRPC3 channel and IP₃R1 primary antibodies, respectively, generated mean N-FRET of ~19% (Fig. 8, B and C). Fig. 8B illustrates that immunoFRET was localized both to the plasma membrane and intracellularly. MβCD treatment and ANT-CSD peptide reduced mean N-FRET to ~12 and 10%, respectively (Fig. 8C). The MβCD-induced reduction in N-FRET was reversed by subsequent exposure of MβCD-treated cells to Chol/MβCD (Fig. 8C). These data indicate that caveolae and the cav-1 CSD maintain IP₃R1 and TRPC3 in close spatial proximity in arterial smooth muscle cells.

**DISCUSSION**

Data in this study indicate that cav-1 maintains IP₃R1 and TRPC3 channels in a signaling complex that enables IP₃-induced Iₐcat activation and vasoconstriction in cerebral artery smooth muscle cells. We show that: 1) caveolae disassembly reduces myogenic tone and attenuates IP₃-induced Iₐcat activation and vasoconstriction, 2) cav-1 knockdown does not alter IP₃R1 and TRPC3 channel expression but reduces myogenic tone and IP₃-induced vasoconstriction, 3) the N and C termini of IP₃R1 and the C terminus of TRPC3 contain CSD-interacting motifs, 4) a CSD peptide inhibits IP₃-induced Iₐcat activation and vasoconstriction, 5) cav-1, IP₃R1 and TRPC3 are located within the same macromolecular complex in cerebral arteries, and 6) the close spatial proximity of IP₃R1 and
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TRPC₃ is disrupted by caveolae disassembly and a CSD peptide. In summary, data indicate that cave-1 and caveolae spatially localize IP₃₃₃₃, R₁ and TRPC₃ channels in cerebral artery smooth muscle cells, enabling IP₃₃₃₃-induced physical coupling, leading to ICₐ₅₁ generation and vasoconstriction.

Three IP₃₃₃₃, R isoforms (1–3) have been cloned (27, 28). IP₃₃₃₃, R₁ is the primary molecular and functional isoform expressed in arterial smooth muscle cells (6). In cerebral artery smooth muscle cells, plasma membrane TRPC₃ channels are located in close spatial proximity to IP₃₃₃₃, R₁ (4). IP₃₃₃₃ activates plasma membrane TRPC₃ channels by inducing a direct interaction between the IP₃₃₃₃, R₁ N terminus and TRPC₃ channel calmodulin and IP₃₃₃₃, R binding (CIRB) domains (4). IP₃₃₃₃-induced TRPC₃ channel activation causes Na⁺ influx, resulting in membrane depolarization, voltage-dependent Ca²⁺ channel activation, an intracellular Ca²⁺ ([Ca²⁺]ᵢ) elevation, and vasoconstriction. Therefore, IP₃₃₃₃, R₁ and TRPC₃ channels directly couple in arterial smooth muscle cells (4, 5).

Several TRP channels, including TRPC₃, localize within caveolae in mammalian cells (29). Such localization may enhance TRP channel interaction with other cell signaling molecules (29). However, whether caveolae localize TRPC₃ channels nearby other regulatory proteins in arterial smooth muscle cells was unclear. Our data indicate that cave-1 and TRPC₃ are spatially localized in the arterial smooth muscle cell plasma membrane, suggesting that TRPC₃ may be localized in caveolae. In rat tail arteries and pulmonary artery smooth muscle cells, cholesterol depletion by MβCD reduced caveolae density and altered plasma membrane morphology, and these effects were reversed by cholesterol replenishment (15, 23). To examine a role for caveolae and cave-1 in enabling functional IP₃₃₃₃, R₁ to TRPC₃ channel physical coupling in arterial smooth muscle cells, we used MβCD at concentrations previously utilized to effectively disrupt plasma membrane caveolae (15, 23, 30, 31). MβCD reduced smooth muscle cell membrane capacitance and this was reversed by subsequent cholesterol treatment. Reduced cell capacitance has also been reported in arterial smooth muscle cells lacking cave-1, and in MβCD-treated uterine and urinary bladder smooth muscle cells (13, 32, 33). These studies and ours indicate that acute and chronic loss of caveolae reduces smooth muscle cell surface area. In another previous study, MβCD-mediated caveolae disruption did not alter arterial smooth muscle cell membrane capacitance (34). Reasons for these contradictory observations are unclear. MβCD modulates depolarization of skinned skeletal muscle fibers (35). However, MβCD treatment did not alter membrane potential or basal intracellular Ca²⁺ concentration in cerebral artery smooth muscle cells (34). These studies suggest that the effect of MβCD on membrane potential may be cell type-dependent. Here, MβCD and cave-1 knockdown attenuated myogenic constriction in cerebral arteries, consistent with observations in cerebral and mesenteric arteries from cave-1 knock-out mice and MβCD-treated rat mesenteric arteries (16, 17, 30). It is unclear why MβCD and cave-1 knockdown reduced myogenic tone, whereas the CSD peptide did not alter myogenic tone. One explanation may be that cave-1 regulates myogenic tone via a CSD-independent mechanism. Alternatively, the CSD peptide used may more effectively block physical coupling of IP₃₃₃₃, R₁ to TRPC₃ channels than it inhibits CSD-dependent mechanisms that lead to cerebral artery myogenic constriction.

We demonstrate for the first time that caveolae disassembly and a CSD peptide reduces IP₃₃₃₃-induced ICₐ₅₁ activation in smooth muscle cells and vasoconstriction in pressurized arteries. Co-IP data demonstrated that cave-1, IP₃₃₃₃, R₁, and TRPC₃ channels are contained within the same macromolecular complex in cerebral arteries. We also show that both IP₃₃₃₃, R₁ and TRPC₃ channels contain CSD binding motifs. The CSD peptide did not alter smooth muscle cell capacitance, suggesting that caveolae integrity and thus, cell surface area remained unchanged. However, the CSD peptide attenuated IP₃₃₃₃, induced ICₐ₅₁ activation in smooth muscle cells and vasoconstriction in pressurized arteries. We tested the hypothesis that caveolae/cave-1 spatially localize plasma membrane TRPC₃ channels nearby SR membrane IP₃₃₃₃, R₁ channels, thereby permitting physical coupling of these proteins. The Förster distance between Alexa 488 and 546 is 6.4 nm (Invitrogen). Given that secondary antibodies bound to IP₃₃₃₃, R₁ (Alexa 546) and TRPC₃ (Alexa 488) generated significant N-FRET, these data indicate that IP₃₃₃₃, R₁ and TRPC₃ channels are located in very close spatial proximity. Our data indicate that FRET was localized both at the plasma membrane and within the cytosol. Intracellular protein complexes containing both IP₃₃₃₃, R₁ and TRPC₃ have previously been reported in HEK-293 cells (36). We have also previously shown that a proportion of TRPC₃ protein is intracellular in rat cerebral arteries and that TRPC₃ immunofluorescence exhibits both plasma membrane and intracellular localization in arterial smooth muscle cells (4). Conceivably, intracellular TRPC₃ channels may be located on organelle membranes that are nearby SR IP₃₃₃₃, R₁ channels or stored within the SR prior to trafficking and in close proximity to SR IP₃₃₃₃, R₁. Caveolae disruption and the CSD peptide both reduced the IP₃₃₃₃, R₁/TRPC₃ N-FRET signal, indicating that these proteins became spatially separated. These data indicate that cave-1, IP₃₃₃₃, R₁ and TRPC₃ channels form a macromolecular complex. Additional data indicate that this complex enables physically coupling of SR IP₃₃₃₃, R₁ to plasma membrane TRPC₃ channels thereby allowing direct communication between intracellular and plasma membrane proteins. MβCD and CSD peptide reduced IP₃₃₃₃, induced ICₐ₅₁ and MβCD, CSD peptide, and cave-1 knockdown attenuated IP₃₃₃₃, induced vasoconstriction. Therefore, cave-1 and caveolae enable IP₃₃₃₃, R₁ to physically couple to TRPC₃ channels in smooth muscle cells to control arterial diameter.

Recombinant cave-1 interacts with both endogenous IP₃₃₃₃, R₃ and recombinant TRPC₁ in HEK-293 cells (37). Expression of a CSD-deleted cave-1 mutant in HEK-293 cells and a human dermal microvascular endothelial cell line increased store-operated Ca²⁺ entry (SOCE) and attenuated a thapsigargin-induced IP₃₃₃₃, R₃-TRPC₁ interaction in HEK-293 cells (37). These studies suggested that through its scaffolding domain, cave-1 regulates an IP₃₃₃₃, R₃-TRPC₁ complex and SOCE in cultured HEK-293 and endothelial cells (37, 38). Store-operated cation currents have been described in a variety of vascular smooth muscle cells, including aorta, mesenteric artery, and portal vein (39). In contrast, SR Ca²⁺ depletion does not acti-
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deplete a I$_{Ca}$ or alter IP$_3$-induced I$_{Ca}$ activation in cerebral artery smooth muscle cells or modify IP$_3$- and agonist-induced cerebral artery constriction at low intravascular pressure (5, 6). SR Ca$^{2+}$ depletion also did not induce contraction in cerebral arteries at low or no pressure (5, 40, 41). Collectively, these studies indicate that SOCE does not appear to exist in contractile cerebral smooth muscle cells.

In conclusion, we demonstrate that cav-1 maintains IP$_3$R1 and TRPC3 channels in a functional macromolecular complex in cerebral artery smooth muscle cells. Cav-1 enables IP$_3$ to stimulate physical coupling between IP$_3$R1 and TRPC3 channels, leading to I$_{Ca}$ activation and vasoconstriction.

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