Transient Receptor Potential Canonical Channels Are Required for *in Vitro* Endothelial Tube Formation*^S

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Background: Different Ca^{2+} entry pathways co-exist in endothelial cells.

Results: On artificial basement membrane, endothelial cells displayed Ca^{2+} oscillations and formed tubes, both processes prevented after silencing TRPC channels, but not Orai1.

Conclusion: TRPC channels are essential for *in vitro* tubulogenesis.

Significance: TRPC channels represent interesting candidates to modulate angiogenesis.

In endothelial cells Ca²⁺ entry is an essential component of the Ca²⁺ signal that takes place during processes such as cell proliferation or angiogenesis. Ca²⁺ influx occurs via the storeoperated Ca²⁺ entry pathway, involving stromal interaction molecule-1 (STIM1) and Orai1, but also through channels gated by second messengers like the transient receptor potential canonical (TRPC) channels. The human umbilical vein-derived endothelial cell line EA.hy926 expressed STIM1 and Orai1 as well as several TRPC channels. By invalidating each of these molecules, we showed that TRPC3, TRPC4, and TRPC5 are essential for the formation of tubular structures observed after EA.hy926 cells were plated on Matrigel. On the contrary, the silencing of STIM1 or Orai1 did not prevent tubulogenesis. Soon after being plated on Matrigel, the cells displayed spontaneous Ca²⁺ oscillations that were strongly reduced by treatment with siRNA against TRPC3, TRPC4, or TRPC5, but not siRNA against STIM1 or Orai1. Furthermore, we showed that cell proliferation was reduced upon siRNA treatment against TRPC3, TRPC5, and Orai1 channels, whereas the knockdown of STIM1 had no effect. On primary human umbilical vein endothelial cells, TRPC1, TRPC4, and STIM1 are involved in tube formation, whereas Orai1 has no effect. These data showed that TRPC channels are essential for in vitro tubulogenesis, both on endothelial cell line and on primary endothelial cells.

In response to circulating agents and/or physical forces, endothelial cells synthesize and release several compounds that act on the underlying smooth muscle cells to modulate their contractile status. In addition, endothelial cells control the regulation of vascular permeability, blood coagulation, or the formation of new blood vessels. Elevation of the cytosolic Ca²⁺ concentration due to Ca²⁺ release from the endoplasmic reticulum (ER),² and Ca²⁺ influx is pivotal for all these functions (1).



In nonexcitable cells, Ca²⁺ can enter via the well described store-operated Ca²⁺ entry (SOCE) pathway and/or through other routes that do not depend on the filling state of the ER. These routes are linked to the generation of second messengers produced upon agonist stimulation, that we called collectively receptor-activated Ca^{2+} entry, RACE (2, 3). SOCE was initially described in 1986 by Putney (4) and linked the level of ER Ca^{2+} store depletion with the opening of plasma membrane Ca²⁺ channels. Recently, two major SOCE components have been identified: the stromal interacting molecule-1 (STIM1) (5-7), located predominantly in the ER membrane and which senses the luminal Ca²⁺ concentration, and the Ca²⁺ channel Orai1 (8-10), which allows Ca²⁺ entry. STIM1 oligomerizes and forms punctae upon store depletion and translocates close to the plasma membrane where it binds to and activates Orai1 (11). In addition to STIM1 and Orai1, channels from the TRP (transient receptor potential) family were shown to constitute an alternative and/or additional Ca^{2+} influx pathway in several cellular systems. The TRP channels (12) are composed of 28 members that are divided into six subfamilies. Among them the TRPC (C for canonical) subfamily comprises both store-operated and non-store-operated cation channels (13). Endothelial cells express a great variety of TRP channels, and more and more data link TRP channel isoforms to specific endothelial cell functions, such as control of the vascular tone, vascular permeability, or angiogenesis (14). Angiogenesis, the formation of new blood vessels from preexisting ones, is a complex and multistep process that comprises migration, proliferation, and reorganization of endothelial cells (15). Ca^{2+} signals are pivotal for many steps that are taking place during angiogenesis (16), and recent data reported that TRPC6 is an important player in angiogenesis (17, 18) whereas others revealed that angiogenesis is severely impaired after Orai1 knockdown (19). In a recent paper, we showed that TRPC1, TRPC3, TRPC4, TRPC5, and TRPC6 isoforms are expressed, together with STIM1 and Orai1, on EA.hy926 cells (20), an endothelial cell line derived from human umbilical vein endothelial cells (HUVECs) fused with human lung adenocarcinoma cell line A549 (21). In this cell line, massive store depletion achieved after thapsigargin stimulation induced Ca²⁺ entry that is mediated both by store-dependent STIM1/Orai1 and store-independent pathways involving TRPC3 (20). In the present study, we focused on

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This article contains supplemental Table 1, Figs. S1–S5, and Movies 1 and 2.
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² The abbreviations used are: ER, endoplasmic reticulum; SOCE, store-operated Ca²⁺ entry; STIM1, stromal interaction molecule-1; STIM2, stromal interaction molecule-2; TRP, transient receptor potential; TRPC, TRP canonical.

the role played by the different Ca^{2+} entry channels in angiogenesis, an important physiological but also pathological process involving endothelial cells, using an *in vitro* approach. We show that TRPC3, TRPC4, and TRPC5 isoforms play a critical role in *in vitro* tube formation, as cells depleted of these proteins were not able to form tubular structures when plated on the artificial basement membrane Matrigel. The absence of tube formation was correlated with a marked decreased of spontaneous Ca^{2+} oscillations displayed by control cells seeded on the extracellular matrix. Once the cells have formed tubes, the Ca^{2+} response elicited by histamine displayed oscillations that depend on the same three TRPC channels. Finally, we showed that on primary HUVEC, TRPC1, TRPC4, and STIM1, but not Orai1, are required for tubulogenesis.

EXPERIMENTAL PROCEDURES

Materials—DMEM, penicillin, and streptomycin were obtained from Invitrogen. Fetal calf serum (FCS) was from PPA Laboratories (Linz, Austria). Histamine, SK&F96365, Pyr3, U73122, and thapsigargin were obtained from Sigma. The acetoxylmethyl ester form of Fura-2 (Fura-2/AM) and BAPTA (BAPTA/AM) were from Molecular Probes Europe (Leiden, The Netherlands). BrdU was from Calbiochem, and the growth factor-reduced Matrigel was from BD Biosciences. The ER-targeted cameleon probe $D1_{ER}$ was kindly provided by Drs. Amy Palmer and Roger Tsien.

Cell Culture and Transfection—Experiments were performed on the HUVEC-derived cell line EA.hy926 (kindly provided by Dr. C. J. S. Edgell) at passages >45. Cells were grown in DMEM containing 15% FCS, 1% HAT (5 mM hypoxanthin, 20 μ M aminopterin, 0.8 mM thymidine), 50 units/ml penicillin, 50 μ g/ml streptomycin and were maintained at 37 °C in 5% CO₂ atmosphere. Primary HUVECs were grown in EGM-2 supplemented with 2% fetal bovine serum (FBS) and Bulletkit (Lonza, Switzerland). For the D1_{ER} (generously provided by Drs. A. Palmer and R. Tsien) experiments, cells were grown until 70–80% confluence and were transiently transfected with 2 μ g of cDNA encoding the D1_{ER} construct.

siRNA Knockdown—EA.hy926 and HUVECs were transfected in suspension by incubating 3×10^5 cells in a solution containing 6 μ l of Lipofectamine RNAiMax (Invitrogen) and the specific siRNA (100 nm; Ambion, Invitrogen, or Qiagen) according to manufacturer's protocols (Invitrogen). The optimal effect is obtained 48 h after transfection. The siRNA scramble from Ambion was used as a negative control. All siRNA used are the same as published previously (20). The other specific siRNA against TRPC3, TRPC4, and TRPC5 used in supplemental Fig. S3 are described in supplemental Table 1.

Cytosolic Calcium Measurements—For Ca²⁺ imaging, EA.hy926 and HUVECs were plated on 30-mm glass coverslips. The changes in cytosolic Ca²⁺ concentration were measured with Fura-2. Cells were loaded with 2 μ M Fura-2/AM plus 1 μ M pluronic acid for 30 min in the dark at room temperature in a medium containing 135 mM NaCl, 5 mM KCl, 2 mM CaCl₂, 1 mM MgCl₂, 10 mM Hepes, 2.6 mM NaHCO₃, 0.44 mM KH₂PO₄, 10 mM glucose with 0.1% vitamins and 0.2% amino acids, pH adjusted to 7.45 with NaOH. Cells were washed twice and equilibrated for 10–15 min in the same buffer to allow deesterifica-



FIGURE 1. **Ca²⁺ dependence of the** *in vitro* **tube formation process on EA.hy926 cells.** The EA.hy926 were seeded on growth factor-reduced Matrigel and visualized 10–14 h later, in control conditions, or in the presence of 30 μ M SK&F96365 (a Ca²⁺ entry blocker), or 50 μ M BAPTA/AM (an intracellular Ca²⁺ chelator). Bar graph shows the percent of the surface covered by tubes in control condition and upon SK&F96365 or BAPTA/AM treatment. Data are mean \pm S.E. (*error bars*) of three independent experiments. *Scale bars*, 200 μ m.***, p < 0.001.

tion. Ratiometric images of Ca^{2+} signals were obtained using a microscope (Axio Observer, Zeiss) equipped with a Lambda DG4 illumination system (Sutter Instrument Company, Novato, CA), which rapidly changed the excitation wavelengths between 340 nm (340AF15; Omega Optical) and 380 nm (380AF15; Omega Optical). Emission was collected through a 415DCLP dichroic mirror and a 510WB40 filter (Omega Optical), by a cooled, 12-bit CCD camera (CoolSnap HQ; Ropper Scientific, Trenton, NJ). Image acquisition and analysis were performed with the Metafluor 6.3 software (Universal Imaging, West Chester, PA, USA). Experiments were performed at room temperature in Hepes-buffered solution containing 135 mM NaCl, 5 mM KCl, 1 mM MgCl₂, 2 mM CaCl₂, 10 mM Hepes, 10 mM glucose, pH adjusted at 7.45 with NaOH. The Ca²⁺-free solution contained 1 mM EGTA instead of 2 mM CaCl₂.

Spontaneous Ca^{2+} signals of cells plated on Matrigel were recorded during 10 min, in culture medium without phenol red at 37 °C. The signal was analyzed as the percentage of responding cells, and the number of Ca^{2+} transients recorded during 10 min, considered only for cells that responded at least with one oscillation.

ER Ca^{2+} *Measurements*—EA.hy926 cells were transiently transfected with 2 μ g of cDNA encoding the D1_{ER} construct 48 h before the experiments. Cells were illuminated at 440 nm (440AF21; Omega Optical), and emission was collected through a 455DRLP dichroic mirror, alternatively at 480 nm (480AF30; Omega Optical) and 535 nm (535AF26; Omega Optical). When required, photobleaching was corrected and data were expressed as R/R_o , R_o being the mean ratio values obtained during the first minute of the recording.



FIGURE 2. Effects of protein silencing on tube formation on EA.hy926 cells. *A*, 48 h after transfection with siRNA (against STIM1, Orai1, TRPC3, TRPC4, and TRPC5), the EA.hy926 cells were seeded on growth factor-reduced Matrigel and visualized 10–14 h later. *B*, tube formation was quantified in control condition and upon siRNA treatment. Data are mean ± S.E. (*error bars*) of three to five independent experiments. *, *p* < 0.05; ***, *p* < 0.001. *C*, cells were seeded on growth factor-reduced Matrigel in control conditions and in the presence of 10 µM TRPC3 inhibitor Pyr3 or 2 µM phospholipase C inhibitor U73122. Scale bars, 200 µm.

Western Blotting—Western blotting was performed as described previously (20). Briefly, the endothelial cells were lysed using modified Nonidet P-40 cell lysis buffer (Invitrogen), 50 mM Tris, pH 7.4, 250 mM NaCl, 5 mM EDTA, 50 mM NaF, 1 mM Na₃VO₄, and 1% Nonidet P-40. Total proteins were separated on SDS-PAGE and transferred to nitrocellulose membranes. Membranes were incubated in T-TBS (0.1% Tween 20, 20 mM Tris-HCl, pH 7.5, 137 mM NaCl) and 5% nonfat milk. Blots were incubated with the primary antibodies diluted in T-TBS and nonfat milk as follows: rabbit anti-TRPC1 polyclonal antibody (1:200; Alomone), rabbit anti-TRPC4 polyclonal antibody (1:200; Alomone), rabbit anti-TRPC5 polyclonal antibody (1:200; Alomone), rabbit anti-TRPC6 polyclonal antibody (1:200; Alomone), and mouse monoclonal antibody against

 α -tubulin (clone DM1A, Sigma) 1:10,000. Blots were incubated with horseradish peroxidase (HRP)-conjugated goat antimouse diluted 1:10,000 (Bio-Rad) or with HRP-conjugated goat anti-rabbit diluted 1:10,000 (Bio-Rad), respectively. Antibodies were revealed using ECL reagents and hyperfilm ECL (Amersham Biosciences). ImageJ software was used to quantify the level of protein expression.

Quantitative Real-time PCR—Real-time PCR experiments were performed at the genomics platform of the NCCR Frontiers in Genetics (Geneva). For each PCR, 1/10 of the cDNA template was PCR-amplified in a 7900HT SDS System using Power SYBR Green PCR master mix (both from Applied Biosystems, Foster City, CA). We obtained raw threshold cycle (Ct) values using SDS 2.0 software (Applied Biosytems). A mean quantity was calculated from triplicate PCRs for each sample,



and this quantity was normalized to the average of three endogenous control genes (glucuronidase B, GAPDH, and TBP) as described in Ref. 22. Primers used for the real-time PCR are described in Ref. 20.

Cell Proliferation Assay—EA.hy926 cells were cultured in 96-well plates at a density of 10000 cells/well. The Calbiochem BrdU cell proliferation assay is a nonisotopic immunoassay for the quantification of BrdU incorporation into newly synthesized DNA of actively proliferating cells. BrdU reagent was added 48 h after cell transfection, for 24 h. Absorbance was measured at 450 nm in each well using a spectrophotometric plate reader, and background was corrected by removing absorbance measured at 590 nm.

Morphogenesis Assay on Matrigel—24-well plates were precoated with a growth factor-reduced Matrigel, and 48 h after transfection EA.hy926 cells (4.5×10^5 cells/well) or HUVECs (1×10^5 cells/well) were seeded onto the Matrigel surface. Cells were incubated for 12–14 h at 37 °C, in the normal culture medium with serum but without addition of extra growth factors, and tubular network visualization was performed using ImageXpress Micro (MDS Analytical Technologies) automated fluorescence imaging system. Tube formation was quantified as the percentage of the cell surface covered by tubular structures.

Statistics—Results are expressed as mean \pm S.E. of *n* observations. Sets of data were compared with a Student's *t* test. Differences were considered statistically significant when *p* < 0.05. *ns*, nonsignificant difference; *, *p* < 0.05; **, *p* < 0.01; ***, *p* < 0.001. All statistical tests were performed using GraphPad Prism version 5.0 for Windows (GraphPad Software).

RESULTS

Role of Ca^{2+} Entry Channels for in Vitro Tube Formation on EA.hy926 Cells—To study the role of Ca²⁺ entry channels in tubulogenesis, EA.hy926 cells were subjected to an in vitro tube formation assay. When endothelial cells were grown on a reconstituted basement membrane matrix Matrigel (23), they spontaneously formed tubular structures in the control condition (Fig. 1). The quantification of this process was performed by measuring the percentage of the surface covered by tubular structures in different conditions. We first evaluated the importance of Ca^{2+} signaling and Ca^{2+} entry in the process of tube formation. As expected, 50 µM BAPTA-AM (an intracellular Ca²⁺ chelator) completely abolished tube formation. Moreover, the nonspecific Ca^{2+} entry blocker SK&F96365 (30 μ M), decreased tube formation by 70% (Fig. 1). These results confirmed the essential role of Ca^{2+} entry/ Ca^{2+} signaling in the *in* vitro tube formation process.

EA.hy926 expressed TRPC1, TRPC3, TRPC4, TRPC5, and TRPC6 together with STIM1 and Orai1 (20). We thus knocked down each of these proteins involved in Ca²⁺ entry and quantified the impact on tube formation. Knockdown of STIM1 or Orai1 proteins evoked a small but significant increase in the surface covered by tubes by approximately 30% (Fig. 2, *A* and *B*). On the contrary, the knockdown of TRPC3 reduced the tube formation by 80% (Fig. 2, *A* and *B*). The pharmacological TRPC3 inhibitor PyR3 (24) completely abolished tube formation, confirming the importance of TRPC3 for tube formation (Fig. 2*C*). TRPC4 and TRPC5 were also important for the *in*



FIGURE 3. **EA.hy926 cell proliferation assay.** *A*, quantification of BrdU incorporation in control conditions and in the presence of 30 μ M SK&F96365 and 50 μ M BAPTA/AM. *B*, quantification of BrdU incorporation in EA.hy926 cells transfected with control siRNA and siRNA against STIM1, Orai1, and STIM1/ Orai1. *C*, quantification of BrdU incorporation in EA.hy926 cells transfected with control siRNA and siRNA against TRPC3, TRPC4, and TRPC5. Data are mean \pm S.E. (*error bars*) of minimum three independent experiments in duplicate. *ns*, not significant; ***, *p* < 0.001.

vitro tube formation process, as the silencing of these proteins reduced tube formation by 65 and 50%, respectively. On the other hand, TRPC1 silencing had no significant impact on this process, and TRPC6 knockdown marginally reduced tube formation (supplemental Fig. 1*A*). We previously reported that STIM1, Orai1, and TRPC3 silencing decreased protein levels by 67-80% (20), and siRNA against TRPC4 and TRPC5 also reduced protein levels by 65 and 70%, respectively (supplemental Fig. 2, *A* and *B*,). We verified, at the mRNA level, the absence of cross-effect among TRPC3, TRPC4, and TRPC5 following their respective knockdown (supplemental Fig. 3). We obtained similar effects on tube formation using different siRNA against TRPC3, TRPC4, and TRPC5 (supplemental Fig. 4*B*).

TRPC channels are known to be activated downstream from the phospholipase C pathway (12). Consequently, we applied the general phospholipase C inhibitor U73122 during the process of tube formation. As shown on Fig. 2*C*, this treatment completely abolished Matrigel-induced tube formation.

Role of Ca^{2+} Entry Channels in EA.hy926 Cell Proliferation— In vivo, angiogenesis requires cell proliferation, in addition to cell migration and organization of tubular structures. Endothelial cell proliferation was evaluated by measuring BrdU incorporation for 24 h. First, we incubated the cells with 50 μ M BAPTA-AM to prevent Ca²⁺ elevations, or with 30 μ M





FIGURE 4. **Spontaneous Ca²⁺ oscillations on EA.hy926 cells during tube formation.** A-D, cytosolic Ca²⁺ measured at 37 °C on EA.hy926 cells loaded with 2 μ M Fura-2 (supplemental Movies 1 and 2). Intracellular Ca²⁺ variations were measured after the first 1–2 h of tube formation on Matrigel, after siControl treatment (A), siSTIM1 (B), siTRPC3 (C) or siTRPC4 (D). Acquisition was done for 10 min, and each *panel* displays the responses of a representative coverslip. E, number of Ca²⁺ oscillations during 10 min in cells treated with siControl, siSTIM1, siOrai1, siTRPC3, siTRPC4, and siTRPC5. F, percentage of cells showing at least one Ca²⁺ transient during the 10-min recording time (n ranges from 64 to 168 cells). *Error bars*, S.E. ns, not significant; **, p < 0.01; ***, p < 0.001.

SK&F96365 to prevent Ca²⁺ entry. In both conditions, we observed a decreased in cell proliferation by approximately 50% (Fig. 3*A*). Then, we estimated the involvement of STIM1/Orai and the different TRPC channels in endothelial cell proliferation. The cell proliferation was not affected by STIM1 knockdown, whereas the silencing of Orai1 decreased cell proliferation by 70% (Fig. 3*B*). The double knockdown (Orai1/STIM1) reduced the proliferation to the same level as siRNA against Orai1 alone (Fig. 3*B*). TRPC3 and TRPC5 knockdown decreased cell proliferation by 65%, whereas the silencing of TRPC4 had no significant impact (Fig. 3*C*). TRPC1 or TRPC6 silencing did not alter endothelial cell proliferation (supplemental Fig. 1*B*).

Spontaneous Ca^{2+} Transients during the Early Phase of in Vitro Tubulogenesis on EA.hy926 Cells—During the first hours after plating on Matrigel, we observed spontaneous Ca^{2+} oscillations in about 55% of the cells (Fig. 4, *A*, *E*, and *F*). To quantify the impact of gene silencing on these Ca^{2+} oscillations, we imaged the cells for 10 min and analyzed the oscillation frequency and the percentage of cells presenting Ca^{2+} transients.

In STIM1- or Orai1-depleted cells, the frequency and the percentage of responding cells were not significantly modified. In TRPC3, TRPC4, or TRPC5 knockdown cells, the spontaneous Ca^{2+} pattern was strongly affected (Fig. 4, *C*–*F*), as shown by the decrease in the number of oscillations (Fig. 4*E*). The percentage of cell oscillating was decreased by 50% in TRPC3- and TRPC5-depleted cells, whereas siRNA against TRPC4 did not significantly affect this parameter (Fig. 4*F*). In line with the effect observed on tube formation, the silencing of TRPC1 or TRPC6 did not alter the spontaneous Ca^{2+} signal (supplemental Fig. 5, *A* and *B*). Similar effects were obtained using the other siRNA against TRPC3, TRPC4, and TRPC5 (supplemental Fig. 4, *C* and *D*). Hence, the alteration of spontaneous Ca^{2+} signals after gene silencing correlated well with the effect observed on tube formation.

Histamine Induced Ca^{2+} Oscillations in EA.hy926 Cells Forming Tubular Structures—10–14 h after adhesion on Matrigel, once the tubular structures are formed, the cells did not display spontaneous Ca^{2+} transients any more (data not shown). Hence, we verified whether the cells were still able to respond to an agonist like histamine. Indeed, the addition of 1





FIGURE 5. **Histamine-induced Ca²⁺ response on EA.hy926 cells after tube formation.** Cytosolic Ca²⁺ responses were measured at 37 °C once the tubular structures were formed (14 h after cell adhesion on Matrigel). Before stimulation, no Ca²⁺ oscillations were detected. *A–D*, Ca²⁺ responses elicited by histamine after siControl treatment (*A*), siSTIM1 (*B*), siOrai1 (*C*), or siTRPC3 (*D*). *E*, number of Ca²⁺ oscillations during 10 min in cells treated with siControl, siSTIM1, siOrai1, siTRPC3, siTRPC4, and siTRPC5. *F*, amplitude of Ca²⁺ oscillations for each conditions shown in *E* (n ranges between 25 and 91 cells). *Error bars*, S.E. *ns*, not significant; **, p < 0.01; ***, p < 0.001.

 μ M histamine induced regular Ca²⁺ oscillations in control cells (Fig. 5*A*). We thus wanted to explore the molecules important for this Ca²⁺ response, even though tube formation was not affected by the presence of histamine (data not shown). The Ca²⁺ oscillatory phenotype was not significantly affected after STIM1 knockdown (Fig. 5, *B*, *E*, and *F*). On the contrary, after Orai1, TRPC3, TRPC4, or TRPC5 silencing, the frequency of Ca²⁺ oscillations was decreased by 32, 80, 40, and 91%, respectively (Fig. 5*E*). The amplitude of Ca²⁺ oscillations was not affected by Orai1 knockdown, whereas the silencing of TRPC3, TRPC4, or TRPC5 channels decreased the average amplitude of Ca²⁺ oscillations by 50, 20, and 20%, respectively (Fig. 5*F*). In TRPC1 and TRPC6 knockdown cells the response to histamine was similar to the one obtained in control cells (supplemental Fig. 5, *C* and *D*). The percentage of cells responding to hista-

mine was significantly reduced after TRPC3 and TRPC5 knockdown (data not shown).

When cells were seeded on glass instead than on Matrigel, they formed a monolayer and displayed a different pattern of Ca^{2+} response to histamine. Indeed, 1 μ M histamine induced a rapid increase of the cytosolic Ca^{2+} concentration, which was followed by a sustained plateau during the whole stimulation. The silencing of STIM1 and Orai1 decreased the plateau phase (estimated after 10 min of stimulation) by 40 and 25%, respectively (Fig. 6, *A* and *B*). Knockdown of TRPC3, TRPC4, and TRPC5 had a stronger effect and reduced histamine-induced Ca^{2+} response by 80, 50, and 75%, respectively (Fig. 6, *C* and *D*). The knockdown of TRPC1 decreased Ca^{2+} entry by about 20% whereas TRPC6 did not appear to play a role in histamineinduced Ca^{2+} entry (supplemental Fig. 1*C*).



FIGURE 6. **Histamine-induced Ca²⁺ entry on EA.hy926 cells grown on glass.** Cytosolic Ca²⁺ concentration was assessed on EA.hy926 cells loaded with 2 μ M Fura-2 and stimulated with 1 μ M histamine in 2 mM Ca²⁺-containing medium. *A*, EA.hy926 cells were transiently transfected with control siRNA (scramble, siControl), or siRNA against STIM1 or Orai1. Each *trace* is the mean of a representative coverslip. *B*, quantification of the effects of siSTIM1 and siOrai1 on the difference between the ratio at 10 min and the basal ratio (*n* ranges from 81 to 201 cells). *C*, representative coverslip. *D*, quantification of the effects of siTRPC3. Each *trace* represents the mean of a representative coverslip. *D*, quantification of the effects of siTRPC3, siTRPC4, and siTRPC5 on the difference between the ratio after 10 min of stimulation and the basal ratio (*n* ranges from 67 to 85 cells). *E*, EA.hy926 cells were transiently transfected with the ER-targeted cameleon probe D1_{ER}. Representative recording of a cell stimulated sequentially with 1 μ M histamine in Ca²⁺-containing medium and in Ca²⁺-free medium is shown. *F*, quantification of ER Ca²⁺ release in EA.hy926 cells stimulated by 1 μ M histamine in Ca²⁺-containing medium (*CA*), Ca²⁺-free medium (*CF*), and by 1 μ M hapsigargin (*Tg*) (in CA). *Error bars*, S.E. ***, *p* < 0.001.

Because STIM1 and Orai1 are implicated in histamine-induced Ca²⁺ entry, we investigated whether this low dose of agonist leads to store depletion. To this end, we transfected the EA.hy926 cells with a genetically encoded Ca²⁺ probe targeted to the ER (D1_{ER}). We hardly could measure a decrease of the ER Ca²⁺ concentration upon 1 μ M histamine stimulation in Ca²⁺ containing medium (Fig. 6, *E* and *F*), which is in line with the small ER depletion obtained with 100 μ M histamine (3). In Ca²⁺-free medium, 1 μ M histamine strongly depleted the store to a level similar to thapsigargin (Fig. 6, *E* and *F*), pointing to an efficient cycling of Ca²⁺ from and to the ER when extracellular Ca^{2+} is available (25, 26). Hence, a low dose of histamine activated STIM1/Orai1 even if the ER Ca^{2+} depletion was almost undetectable with the $D1_{FR}$ (see "Discussion").

Role of Ca^{2+} Entry Channels in Primary Cultured Endothelial Cells—We used the primary cultured HUVECs to assess the role of TRPC channels and STIM1/Orai1 during *in vitro* tube formation and spontaneous Ca²⁺ signals. We first determined that TRPC1 and TRPC4 are expressed on HUVECs (Fig. 7*A*) as well as STIM1 and Orai1. The efficiency of gene silencing was assessed at the mRNA level and ranged between 50 and 85% of mRNA reduction (Fig. 7*B*). The knockdown of STIM1, TRPC1,





FIGURE 7. Expression of TRPC isoforms on HUVEC and effects of protein silencing on tube formation. *A*, mRNA amplification of TRPC1, TRPC3, TRPC4, TRPC5, TRPC6, and TRPC7 isoforms by real-time PCR in HUVECs. *B*, mRNA levels assessed by quantitative RT-PCR 48 h after transfection (mean \pm S.E. (*error bars*), 2–4 independent experiments). *C*, 48 h after transfection with siRNA (against TRPC1, TRPC4, STIM1, and Orai1), the HUVECs were seeded on growth factor-reduced Matrigel and visualized 10–14 h later. *D*, quantification of tube formation in control condition and upon siRNA treatment. Data are mean \pm S.E. of three to five independent experiments. *Scale bar*, 200 μ m. *ns*, not significant; ***, *p* < 0.001.

and TRPC4 reduced tube formation, whereas siRNA against Orai1 had no effect (Fig. 7, *C* and *D*). To ensure that the absence of effect of siOrai1 during tube formation was not due to a moderate efficiency of our siOrai1, we performed Western blotting and found that the protein level of Orai1 was decreased by approximately 80% after siRNA treatment (data not shown). Similar to what we observed on EA.hy926 cells, the HUVECs displayed spontaneous Ca^{2+} oscillations when plated on Matrigel, at the beginning of tubulogenesis (Fig. 8*A*). siRNA against TRPC4 strongly reduced both the frequency and the percentage of responding cells, and the knockdown of TRPC1 and STIM1 reduced Ca²⁺ oscillations to a lesser extend (Fig. 8, B-D). siOrai1 minimally reduced the frequency of Ca²⁺ oscillations (Fig. 8*C*). Hence, on HUVEC (as on EA.hy926 cells), the impact of gene silencing on the spontaneous Ca²⁺ response correlated well with the effect on tube formation.

DISCUSSION

In the present work, we assessed the role of STIM1, Orai1, and TRPC channels on cell proliferation and in vitro tube formation in the HUVEC line, EA.hy926, and on the parental primary cells, the HUVECs. By using a siRNA approach, we showed that on EA.hy926 cells, the knockdown of TRPC3, TRPC4, or TRPC5 strongly decreased Matrigel-induced tube formation, whereas the silencing of STIM1 or Orai1 had no negative impact. Reduction of tube formation was correlated with the decrease of spontaneous Ca^{2+} oscillations during the early stages of the process. We also demonstrated that the silencing of Orai1, TRPC3, or TRPC5 strongly reduced endothelial cell proliferation. On primary cultured endothelial cells (HUVECs), the knockdown of TRPC1, TRPC4, and STIM1 reduced tube formation and spontaneous Ca²⁺ oscillations, whereas the siRNA against Orai1 had no effect on these processes. From these results, it appears that TRPC channels play a major role in tubulogenesis, both in EA.hy926 cells and in HUVECs.

In vitro angiogenesis, using Matrigel as the artificial basement membrane, is widely used to study endothelial cell tube formation (27, 28). In addition, the endothelial cell line EA.hy926 was demonstrated to be a suitable model for *in vitro* studies of tube formation (29-31). Using this assay, we revealed an important role of TRPC3, TRPC4, and TRPC5 channels, which, unlike STIM1 and Orai1, are required for tube formation. On the contrary, the ability to form tubular structures was enhanced upon STIM1/Orai1 gene silencing. The role played by these different Ca²⁺ entry channels correlated well with their role in spontaneous Ca²⁺ oscillations recorded during the early phase of tube formation, which strengthen the link between tube formation and Ca2+ entry. An early study reported an increase in cytosolic Ca²⁺ concentration during the initial phase of endothelial cell sprouting on collagen IV surface (32). The Ca²⁺ rise and the cell sprouting were prevented by the nonspecific Ca²⁺ entry blockers carboxyamidotriazole and SK&F96365 (32). Our study confirmed the essential role of Ca²⁺ entry for endothelial tube formation and shows that TRPC channels are important for the early Ca²⁺ signals associated with tubulogenesis.

TRPC channels have been shown to participate in *in vitro* angiogenesis, in particular TRPC6 (17, 18). On EA.hy926 cells, TRPC6 knockdown slightly reduced tube formation, a marginal effect compared with the important effect of TRPC3, TRPC4, and TRPC5 knockdown. TRPC1 was shown to have a prominent role in angiogenesis in zebrafish (33), but in mammals, no further data reported a role of TRPC channels in tube formation (14).

Angiogenesis is a complex process that comprises not only cell migration and organization as tubular structures, but also requires cell proliferation. On Matrigel, the cells do not prolif-





FIGURE 8. **Spontaneous Ca²⁺ oscillations during the tube formation on HUVECs**. *A* and *B*, cytosolic Ca²⁺ measured at 37 °C on HUVECs loaded with 2 μ M Fura-2. Intracellular Ca²⁺ variations were measured after the first 1–2 h of tube formation on Matrigel, after siControl treatment (*A*) and siTRPC4 (*B*). Acquisition was done for 10 min, and each panel displays the responses of a representative coverslip. *C*, percentage of cells showing at least one Ca²⁺ transient during the 10-min recording time (*n* ranges from 39 to 135 cells). *D*, number of Ca²⁺ oscillations during 10 min in cells treated with siControl, siTRPC1, siTRPC4, siSTIM1, and siOrai1. *ns*, not significant; *, *p* < 0.05; **, *p* < 0.001.

erate (27), but migrate and reorganize to form capillary-like structures. Hence, only part of the angiogenic process is recapitulated using this assay. Consequently, a channel that is important for cell proliferation is likely also important for angiogenesis if we consider the full process. The involvement of Orai1 but not STIM1 in cell proliferation is at odds considering the way Orai1 is getting activated, i.e. upon STIM1 oligomerization, recruitment close to the plasma membrane and binding to Orai1 (11, 34). This, however, was already observed in endothelial cells (35) and in HEK293 cells (36). In both studies, STIM2 was involved in cell proliferation and thus was likely responsible for Orai1 activation. In our model, STIM2 silencing did not impact on cell proliferation (data not shown). Alternatively, Orai1 might be activated independently of STIM1, as was recently reported (37). Furthermore, it was also proposed that Orai1 might interact with cyclin-dependent kinase, pointing to a specific role of this channel in cell proliferation (38).

From our data on EA.hy926 cells, it emerged that STIM1 and Orai1, the major contributors of SOCE, are not required for tubulogenesis, whereas several of the TRPC channels appeared to be. The Ca²⁺ oscillatory behavior observed after stimulation with 1 μ M histamine in cells forming tubes is reminiscent of the spontaneous Ca²⁺ transients observed at the beginning of tube formation. Furthermore, both responses relied mainly on TRPC3, TRPC4, and TRPC5 channels, and minimally on STIM1 and Orai1. That likely implies that the ER Ca²⁺ content stayed elevated during Ca²⁺ oscillations and is in line with previous papers showing that Ca²⁺ oscillations do not affect the Ca²⁺ level within the ER (39) and thus do not activate SOCE

(40). Previously, we and other have shown that 100 μ M histamine led to approximately 15% of store depletion (3), and in these conditions, STIM1 and Orai1 accounted for approximately 35–50% of the sustained Ca²⁺ entry (data not shown). In the present study, we showed that $1 \mu M$ histamine induced also a sustained Ca^{2+} response of cells forming monolayers on glass coverslip, and STIM1/Orai1 knockdown reduced Ca²⁺ entry by approximately 20-40%. The level of store depletion after 1 µM histamine was, however, minimal (Fig. 6E). Hence, it appears that the extent of STIM1/Orai1 involvement in cytosolic Ca²⁺ response is not linearly correlated with the level of store depletion we measured with the $D1_{ER}$. That might be explained by restricted ER compartments emptied by histamine but not measured by the $D1_{ER}$. Recently, it was shown by electron microscopy that proteins targeted to the ER through a KDEL sequence were excluded from thin cortical ER structures (41), suggesting that $D1_{ER}$ might also be excluded from these subcompartments. Whether such regions are depleted by low dose of agonists and whether that leads to the activation of STIM1/Orai1 remained an open question. As already mentioned, during Ca²⁺ oscillations, we can reasonably postulate that the actual level of ER Ca²⁺ depletion is extremely limited, and that would explain that STIM1/Orai1 do not participate in Ca²⁺ entry during Ca²⁺ oscillatory response, at least on EA.hy926 cells. The signaling pathway leading to specific TRPC activation remained, however, to be determined.

Recently, it was reported that Orail channel is an important player for both *in vitro* and *in vivo* angiogenesis (19). The *in vitro* experiments were performed on HUVECs, but neither the



impact of other channels like the TRPC nor the role of STIM1 was evaluated. Hence, we wanted to investigate whether TRPC channels are important for angiogenesis in HUVECs. We found out that the knockdown of TRPC1, TRPC4, and STIM1 reduced tube formation, and, like on EA.hy926 cells, also reduced spontaneous Ca²⁺ oscillations observed at the beginning of tubulogenesis. However, and on the contrary to the findings of Li et al. (19), we did not observe a defect of in vitro angiogenesis after knocking down Orai1. The main difference between both studies is the use of vascular endothelial growth factor (VEGF) to enhance tube formation. We did not add VEGF in the medium whereas Li et al. added 2 ng/ml VEGF. The application of VEGF activated STIM1 and Orai1 (19), whereas without this growth factor, the level of Ca²⁺ within the ER likely remained elevated (as we proposed for the EA.hy926 cells) and thus would explain the absence of Orai1 activation. The mechanism leading to the activation of STIM1 that we observed, remains, however, to be elucidated. We should mention that on HUVECs, it was shown that Orai1 is important for cell proliferation (35). Hence, we cannot exclude an implication of Orai1 during in vivo angiogenesis, as in vivo, cell proliferation is part of the process.

In this study, we showed that *in vitro* tubulogenesis relies on a Ca²⁺ signaling pathway that engages mainly TRPC channels. The knockdown of several TRPC channels, both on an endothelial cell line (EA.hy926) as well as on primary endothelial cells (HUVECs), severely prevented spontaneous Ca²⁺ oscillations of cells plated on Matrigel, as well as the subsequent tube formation. Interestingly, Orai1, a major player in Ca²⁺ signaling, does not seem to be essential for tube formation. Instead, TRPC channels are pivotal and represent very interesting candidates to modulate angiogenesis.

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