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Store-Operated Ca\(^{2+}\) Entry Contributes to Piezo1-Induced Ca\(^{2+}\) Increase in Human Endometrial Stem Cells

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Abstract: Endometrial mesenchymal stem cells (eMSCs) are a specific class of stromal cells which have the capability to migrate, develop and differentiate into different types of cells such as adipocytes, osteocytes or chondrocytes. It is this unique plasticity that makes the eMSCs significant for cellular therapy and regenerative medicine. Stem cells choose their way of development by analyzing the extracellular and intracellular signals generated by a mechanical force from the microenvironment. Mechanosensitive channels are part of the cellular toolkit that feels the mechanical environment and can transduce mechanical stimuli to intracellular signaling pathways. Here, we identify previously recorded, mechanosensitive (MS), stretch-activated channels as Piezo1 proteins in the plasma membrane of eMSCs. Piezo1 activity triggered by the channel agonist Yoda1 elicits influx of Ca\(^{2+}\), a known modulator of cytoskeleton reorganization and cell motility. We found that store-operated Ca\(^{2+}\) entry (SOCE) formed by Ca\(^{2+}\)-selective channel ORAI1 and Ca\(^{2+}\) sensors STIM1/STIM2 contributes to Piezo1-induced Ca\(^{2+}\) influx in eMSCs. Particularly, the Yoda1-induced increase in intracellular Ca\(^{2+}\) ([Ca\(^{2+}\)]\(_i\)) is partially abolished by 2-APB, a well-known inhibitor of SOCE. Flow cytometry analysis and wound healing assay showed that long-term activation of Piezo1 or SOCE does not have a cytotoxic effect on eMSCs but suppresses their migratory capacity and the rate of cell proliferation. We propose that the Piezo1 and SOCE are both important determinants in [Ca\(^{2+}\)], regulation, which critically affects the migratory activity of eMSCs and, therefore, could influence the regenerative potential of these cells.

Keywords: endometrial mesenchymal stem cells; cell migration; mechanosensitive channels; Piezo1 channels; store-operated Ca\(^{2+}\) channels; Ca\(^{2+}\) influx

1. Introduction

Mesenchymal stem cells derived from a wide range of adult tissues, such as bone marrow, umbilical cord, peripheral blood and adipose tissue, are widely applied to treat various diseases. In the past decades, new sources of adult stem cells, including the placenta [1], fallopian tube [2], amniotic fluid [3,4], amniotic membrane [5] and endometrium [6–8] were established. Special interest was given to the stem cells isolated from female reproductive organs, particularly the endometrium [6–8]. The endometrium is a dynamic tissue involved in the synchronized functions of cellular proliferation, differentiation and menstrual shedding [9]. It constantly sheds its top layer of cells during menstruation, which is later regenerated completely by the basal layer of the endometrial tissue, suggesting the existence of a very high potential stem cell population [10]. Endometrial stem cells (eMSCs) are a heterogeneous cellular population composed mainly of stromal and epithelial cells of the endometrial gland that demonstrate an MSC-specific, fibroblast-like morphology [11]. It was found that eMSCs can be differentiated into various types of cells such as myocytes, cardiomyocytes, osteocytes, adipocytes and neurons [12]. Due to their high availability,
these cells are considered as one of the reliable, non-invasive and very valuable substrates for therapeutic interventions. Therefore, profound knowledge of the molecular mechanisms that control the migration and differentiation of eMSCs is of clinical importance.

The mechanical action on main cellular functions, such as proliferation, migration, gene expression and differentiation, is very important for stem cell transplantation since stem cells are constantly exposed to different mechanical stimuli and forms of physical stimulation, including sound, light, temperature, mechanical force and even gravity [13–16]. At the same time, it is established that ion channels, particularly mechanosensitive channels, are directly or indirectly involved in the transduction of all forms of physical stimulation [17]. Piezo1 and Piezo2 molecules were identified as a novel family of mechanosensitive cation channels by Coste et al. [18]. Further, Piezo channels were shown to participate in a variety of mechano-dependent physiological processes, and they were found to play a crucial role in the transmission of physical stimulation to intracellular signaling pathways [19,20]. The signaling functions of Piezo are mainly mediated by their ability to conduct Ca^{2+} ions, which are known to be crucial, secondary messengers in living cells. There is substantial evidence connecting the handling of intracellular calcium concentration ([Ca^{2+}]_i) to the normal physiology and pathophysiology of endometrial stromal cells. It is established that different Ca^{2+}-permeable ion channels are involved in decidualization and play a key role in embryonic functions [21–24].

The participation of Ca^{2+} channels, TRP channels, cAMP-induced cytosolic Ca^{2+} levels and Ca^{2+}-binding proteins suggests that [Ca^{2+}]_i is tightly connected with uterine function [21,25–28]. However, owing to technical and ethical problems, ion channel investigations have been restricted in human endometrial stem cells isolated from human tissue. Here, we use human eMSCs to address the role of mechanosensitive Piezo1 channels in [Ca^{2+}]_i level regulation in endometrial cells. Given the significant role of the temporal and spatial Ca^{2+} regulation in a variety of main cellular functions, including cell motility, we also examined the role of Piezo1 in endometrial cell migration.

2. Results
2.1. Expression of Mechanosensitive Piezo1 Channels in Human Endometrial Mesenchymal Stem Cells (eMSCs)

Piezo1 channels are found in various tissues and cell lines, including several human mesenchymal stem cells [29,30]. The mechanosensitive channel activity with biophysical properties close to the Piezo family was observed in the plasma membrane of eMSCs in response to membrane stretch (“negative” pressure application, suction [31]). Importantly, the Piezo channel family consists of two members [18], Piezo1 and Piezo2, and, of them, only Piezo1 can be activated by both “negative” or “positive” pressure application, whereas Piezo2 was recently reported to only be sensitive to low-threshold “positive” pressure [32]. Therefore, we hypothesized that Piezo1 proteins are likely molecular correlates of stretch-activated MS channels in eMSCs, and they could provide a physiologically relevant pathway for Ca^{2+} influx and, thereby, affect the [Ca^{2+}]_i and Ca^{2+}-dependent signaling processes. By consistently using PCR analysis and immunofluorescence, we first confirmed the expression of Piezo1 in eMSCs (Figure 1A,B).

To detect the functional activity of Piezo1 channels in the plasma membrane of eMSCs, we stimulated the channels in cell-attached patch-clamp experiments using a selective chemical activator of Piezo1 [33]: a small synthetic molecule Yoda1. No significant background channel activity was observed in the control patches (without Yoda1, n > 100, Supplementary Figure S1). Importantly, the presence of Yoda1 in the patch pipette activated the unitary inward currents (Figure 1C) with biophysical properties similar to those previously reported for stretch-activated currents in eMSCs [31]. Moreover, the application of “negative” pressure in the presence of Yoda1 further increased Yoda1-induced channel activity (Figure 1C, the effect is demonstrated on −20 and −40 mV holding potentials). Taken together, our data strongly support the hypothesis of Piezo1 channel expression and its activation by Yoda1 in human eMSCs.
Figure 1. Presence of Piezo1 in eMSCs. (A) RT-PCR analysis revealed the presence of PIEZO1 mRNA. M—size marker, Line 1—primers specific for PIEZO1 amplified the PCR product of the expected size (360 bp), Line 2—RT-PCR negative control in which reverse transcriptase was omitted. Shown is cropped gel with enhanced contrast. (B) Immunofluorescent staining detected Piezo1 proteins (red channel) in eMSCs. No staining of the cells was observed after incubation of the cells with only fluorescent secondary antibodies (Supplementary Figure S2). Cell nuclei were counterstained with DAPI (blue channel). The scale bar is 50 µm. (C) The single-channel activity of Piezo1 induced by selective chemical Piezo1 agonist Yoda1 (10 µM in the pipette solution). Representative current recordings at different membrane potentials are shown. Holding membrane potentials are indicated near current traces, closed and open states indicate the baseline ("zero" current) and Piezo1 open state, respectively. Note that the application of "negative" pressure (p < 0, suction, indicated by the arrows) further increased the activity of the Yoda1-induced channels. The mean I–V relationship corresponded to single-channel conductance of 23.2 ± 1.3 pS (n = 6).

2.2. Piezo1 Channels as Ca²⁺ Entry Pathway in eMSCs

It is well known that the activation of mechanosensitive, cation-selective channels results in Ca²⁺ entry from the extracellular environment to the cytoplasm of the living cells. Thereby, mechanosensitive channels play a significant role in the regulation of [Ca²⁺]i. Here, we assume that stretch-activated MS channels identified as Piezo1 proteins promote Ca²⁺ influx and affect the [Ca²⁺]i in eMSCs. To test this hypothesis, we loaded cells with Fura-2AM, and Ca²⁺ responses induced by selective chemical Piezo1 activator Yoda1 were estimated using an AxioObserver.Z1 inverted microscope imaging system (see Section 4). In our experiments, a large increase in [Ca²⁺]i was observed after the addition of 10 µM Yoda1 to the extracellular solution. Interestingly, the Yoda1-induced Ca²⁺ response consisted of two distinct components: the first, a fast component of [Ca²⁺]i growth developed immediately and quickly declined (plateau) after the addition of Yoda1 to the external solution. Then, several minutes later, a second increase in [Ca²⁺]i (delayed component), which lasted the length of the [Ca²⁺]i registration, was observed (Figure 2A). A calcium ionophore, ionomycin (at a final concentration of 10 µM), was used as a positive control for the reaction (Supplementary Figure S4). Particularly, the addition of ionomycin further increased [Ca²⁺]i, indicating that the second Yoda1-induced component was not caused by disruption of the cell membrane integrity followed by passive calcium entry. Wash-out of extracellular Ca²⁺ with a Ca²⁺-free solution abrogated the delayed Yoda1-induced increase in [Ca²⁺]i, indicating the extracellular nature of the second calcium entry. In addition, as the experiments lasted up to 30 min, we specifically checked the potential damage of the cells by UV light [34]. No visible morphological changes in the cells exposed to UV radiation were observed (Supplementary Figure S3).
Therefore, using PCR analysis, we investigated the expression of the ORAI1–3 products that correspond to STIM1 transcripts, respectively, which are exclusively present in mammals [36]. The STIM1 protein has another homolog: STIM2. Therefore, using PCR analysis, we investigated the expression of the ORAI1–3 and STIM1 and 2 in endometrial stem cells. PCR primers were designed as indicated in Material and Methods. RT-PCR analysis revealed the 78 bp products, 194 bp products and 247 bp products that correspond to ORAI1, STIM1 and STIM2 transcripts, respectively, which are expressed in eMSCs (Figure 3A), whereas the presence of ORAI2 and 3 mRNA was not detected (Supplementary Figure S5).

Figure 2 indicates that the kinetics of Yoda1-induced Ca^{2+} responses resemble the classical Ca^{2+} responses observed due to Ca^{2+} release from internal Ca^{2+} storage with subsequent store-operated Ca^{2+} entrance (SOCE) via the plasma membrane (see, for example, [35]). Therefore, firstly, we examined the potential Yoda1 participation in Ca^{2+} release from Ca^{2+} stores. Figure 2B demonstrates no [Ca^{2+}]], increase after Yoda1 application to cells in calcium-free solution, suggesting no Ca^{2+} release from intracellular Ca^{2+} stores in response to Piezo1 activation. Notably, the further addition of thapsigargin (TG), a specific, irreversible inhibitor of endoplasmic reticulum (ER) Ca^{2+}-ATPases in the same ion conditions resulted in TG-induced Ca^{2+} release, as was previously documented elsewhere [35]. Figure 2B shows the typical Ca^{2+} increase induced by TG in a Ca^{2+}-free solution and the following store-operated Ca^{2+} entry, indicating the existence of functional Ca^{2+} store release and the following activation of SOCE in eMSCs.

2.3. Expression of ORAI1, STIM1 and STIM2 in eMSC Cells

SOCE is a general Ca^{2+} influx via the plasma membrane which is triggered by G-protein-coupled receptors (GPCRs) activation and the following IP3-mediated depletion of endoplasmic reticulum Ca^{2+} stores. SOCE represents a complex of proteins that includes calcium-selective channel ORAI and the endoplasmic reticulum-associated Ca^{2+} sensor STIM. It is known that the ORAI family has three homologs, ORAI1, 2 and 3, and ORAI3 is exclusively present in mammals [36].
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Figure 3. ORAI1, STIM1 and STIM2 expression in human eMSCs. (A) RT-PCR analysis of ORAI1, STIM1 and STIM2 expression in eMSCs. M—size marker. Line 1—primers specific for ORAI1, STIM1 and STIM2 amplified the PCR products of the expected sizes (78, 124 and 247 bp, correspondingly). Line 2—RT-PCR negative control in which reverse transcriptase was omitted. Shown are cropped gels with enhanced contrast. Original gel is shown in Supplementary Figure S5. The distribution of ORAI1 (B), STIM1 (C) and STIM2 (D). Immunoreactivity in eMSCs was examined with confocal microscopy. No staining of the cells was observed after incubation of the cells with only fluorescent secondary antibodies (Supplementary Figure S2). Scale bar is 50 µm.

To evaluate the ORAI1, STIM1 and STIM2 protein localization and spatial distribution in cells, the specific antibodies and immunofluorescence confocal microscopy were utilized. As presented in Figure 3, the immunofluorescence signal was detected in both cytoplasm and plasma membrane of the cells marked with the antibodies recognizing ORAI1 (Figure 3B), STIM1 (Figure 3C) and STIM2 (Figure 3D).

2.4. Contribution of SOCE in Piezo1-Induced Ca\(^{2+}\) Influx

Preliminary experiments showed that the calcium responses to selective chemical stimulation of mechanosensitive Piezo1 channels kinetically resemble that of store-operated Ca\(^{2+}\) entry activated by TG (Figure 2). Therefore, we hypothesized that SOCE may contribute to the increase in [Ca\(^{2+}\)]\(_i\) induced by Piezo1 channel activation. To verify this hypothesis, a pharmacological SOCE inhibitor, 2-aminoethyl diphenylborinate (2-APB), was used. At the beginning of the experiment, Ca\(^{2+}\) was removed from the external cell medium, which immediately led to a reduction in the level of [Ca\(^{2+}\)]\(_i\). Subsequent application of TG caused a rapid [Ca\(^{2+}\)]\(_i\) growth followed by a rapid decline (Figure 4A). The addition of 2 mM Ca\(^{2+}\) to the external solution led to typical growth of [Ca\(^{2+}\)]\(_i\), as a result
of SOCE activation. It is especially noteworthy that the application of 1 µM 2-APB did not suppress the SOCE-activated Ca^{2+} influx, whereas 5 µM 2-APB completely abolished it (Figure 4A).

![Figure 4. 2-APB suppresses the store-operated Ca^{2+} entry and abrogates the second (delayed) phase of Yoda1-induced Ca^{2+} increase in eMSCs. (A) Shown are typical Ca^{2+} responses from the representative experiment (n = 4) following the 1 µM and 5 µM 2-APB application on eMSCs. Measurements were performed in individual cells with the imaging system. [Ca^{2+}]_{i} is plotted versus time. Each point is a mean ± SD from 17 cells. (B) 2-APB suppresses both SOCE and the delayed phase of Piezo-mediated Ca^{2+} entry in eMSCs. Each point is a mean ± SD from 13 cells.](image)

Therefore, to eliminate the potential contribution of SOCE in Piezo-induced Ca^{2+} influx, 2-APB was used in the concentration of 5 µM. Figure 4B demonstrates that the addition of Yoda1, together with 2-APB, to the extracellular solution does not prevent the occurrence of the first component of [Ca^{2+}]_{i} increase but suppresses the second (delayed) one, suggesting the SOCE contribution in Piezo1-mediated Ca^{2+} increase.

2.5. Role of SOCE and Piezo1 in eMSC Migration and Proliferation

It is well known that MSCs possess a homing capacity, can move into damaged areas and help in tissue regeneration [37–41]; therefore, MSCs represent a significant tool for regenerative medicine [42–45]. Lately, there were several reports that proper control of Ca^{2+} signaling is important for effective cell migration. However, how Ca^{2+} coordinates the necessary factors for effective cell moving, key signaling pathways and signaling molecules is still elusive. Here, we investigate the likelihood of participation of Piezo1- or SOCE-induced Ca^{2+} currents in eMSC migration.

The wound healing assay, together with time-lapse imaging, was used to address this challenge. To avoid some possible, uncontrolled effects caused by cell damage, the wound in the eMSC culture was created by the removal of a silicone insert, as described in Materials and Methods. Figure 5A presents typical images showing the wound areas at the starting time point and 36 h after eMSC treatment with 1 µM of TG and 10 µM Yoda1 (see also Supplementary Video S1 showing the time-lapse process of eMSC wound healing). Wound healing area analysis revealed that both SOCE and Piezo1 activation significantly inhibits the wound healing processes. Particularly, the control wounds were fully closed 36 h after the beginning of the experiments, whereas about 40% of the wounds in the presence of Yoda1 or TG remained uncovered by the cells (Figure 5B, Supplementary Table S4). Interestingly, the cell cultivation in the presence of 5 µM 2-APB also resulted in the suppression of the wound healing process (Figure 5B, Supplementary Table S4). To exclude the potential toxic effect of the compounds, we determined the eMSC viability using PI staining at the end of the cell migration experiments. It was established that the percentage of viable eMSCs in the population did not significantly change even after 48 h of treatment with TG, Yoda1 or 2-APB compared to control cells (Figure 5C, Supplementary Table S4). Thus, the
data demonstrated that the observed effect of the suppression of wound healing processes is not caused by the cytotoxic effects of the reagents.

Figure 5. SOCE and Piezo1 in eMSCs migration. (A) Shown are representative images of the experimental wounds at the beginning (0 h) and the end of the wound healing assay (36 h). The scale bar is 200 μm (B). The eMSC wound healing dynamics in the presence of TG, Yoda1 or 2-APB in the culture media. Wound sizes are shown as mean ± SEM (n = 3–4 for each condition). *—significantly different compared to control, Student’s t-test with Welch’s corrections, p < 0.05. (C) No significant changes in the viability of eMSCs were observed. Shown is the mean percent (±SD, n = 3) of dead cells for each experimental condition. #—not significantly different compared to control, two-sample Student’s t-test (p < 0.05).

Wound healing is a standard behavior of epithelial and endothelial cells, as well as mesenchymal cells, that is firstly activated to increase their motility in order to settle the injured area and then, together with their ability to differentiate and to secrete various biologically active factors and molecules [46], they also demonstrate a high, proliferative rise to ensure the successful completion of the wound repair process [47]. Our data revealed the suppression of the wound healing induced by Yoda1, TG and 2-APB; therefore, we assumed that these reagents potentially affect the rate of cell proliferation, and this may partially underlie the observed decrease in cell motility. Our experiments showed that all reagents suppressed the growth of eMSCs, and the pronounced result was observed after 48 h of cell treatment (Figure 6A). The cell cycle FACS analysis showed the significant eMSCs accumulation in the G0/G1 phase after TG treatment (Figure 6B, see, also, Supplementary Table S5). At this time, the pattern of phase distribution of the cells treated with Yoda1 or 2-APB showed no accumulation of the cells in the particular phase of cell cycle.
endometrial stem cells. We also show that the small-molecule modulator of Piezo1 channels, Yoda1, induces large Ca\(^{2+}\) entry in eMSCs. Importantly, the Yoda1-induced Ca\(^{2+}\) response are similar to those induced by TG in numerous cells, particularly: (i) TG-mediated [Ca\(^{2+}\)]\(_i\) growth caused by store depletion followed by (ii) store-operated Ca\(^{2+}\) entry. However, our experiments demonstrated that the application of Yoda1 in a Ca\(^{2+}\)-free solution has no effect on [Ca\(^{2+}\)]\(_i\), whereas the following addition of TG results in the increase of [Ca\(^{2+}\)]\(_i\). Both TG-activated store depletion and store-operated Ca\(^{2+}\) entry were detected in Ca\(^{2+}\) measurements carried out on eMSCs. We also identified that store-operated Ca\(^{2+}\) entry in eMSCs could be formed by ORAI1/STIM1 or ORAI1/STIM2 proteins. In our experiments, a specific blocker of SOCE, 2-APB, suppressed the TG-induced SOCE. Moreover, 2-APB also blocked the second (delayed) component of Yoda1-induced Ca\(^{2+}\) increase, suggesting the SOCE is participating in mechanosensitive Ca\(^{2+}\) signaling in cells. These findings correlate with the data which show that store-operated Ca\(^{2+}\) entry can participate in Ca\(^{2+}\) influx induced by the other agonists, for example, ATP [30,48].

Aside from this, some scientific reports recently indicated that Ca\(^{2+}\) entry mediated by mechanosensitive channels is regulated by a diversity of major proteins which are located in the cellular membrane or endoplasmic reticulum. It was shown that mechanosensitive channels can be physically connected with some Ca\(^{2+}\) regulatory proteins [49]. For example, Piezo and SERCA were discovered to cross-talk to each other through physical interaction in endothelial cells. SERCA2 was demonstrated to suppress Piezo1-induced HUVEC migration via a 14-residue linker region. The linker mutations disrupt the cooperation between Piezo1 and SERCA2, and, consequently, the mechanical activation of Piezo1.
channels is essentially reduced [50]. The main function of sarco/endoplasmic reticulum Ca\(^{2+}\)-ATPase (SERCA) is to transport Ca\(^{2+}\) from the cytosol into the sarco/endoplasmic reticulum. SERCA inhibition induces the store-operated Ca\(^{2+}\) influx. One could, therefore, assume that Piezo1, by closely interacting with SERCA2, can activate SOCE that contributes to the Piezo1-induced Ca\(^{2+}\) entry.

Our results show that TG decreases the migration of eMSCs. These data are consistent with those presented by the other authors that show that TG decreases the proliferation, migration and invasion of human adenocortical carcinoma SW-13, NCI-H295R [51] and esophageal carcinoma EC109 and TE12 cell lines [52]. TG also suppressed ACC xenograft tumor progression in vivo [51]. TG is considered as a potential antineoplastic drug; however, it is also highly toxic to normal cells [53]. Interestingly, we did not observe any effect of TG on eMSC viability even after 48 h of incubation (Figure 5C, Supplementary Table S4).

In addition, we demonstrated that selective chemical activation of Piezo1 by Yoda1 decreases the migration potential of eMSCs. In our previous study, we showed that Yoda1 dose-dependently inhibits migration of transformed mouse fibroblasts [54] that was accompanied by F-actin assembly and stress fiber formation. It is noteworthy that the effect of selective Piezo1 activation by Yoda1 on cellular motility remains rather controversial. Particularly, the increase of migration of mesenchymal stem cells in response to Yoda1 was reported, and the effect was mediated via the release of ATP and the activation of P2 receptor signaling [30]. In contrast, the decrease of migration of keratinocytes in response to Yoda1 was observed, and the specific mechanism which is the localized increase of the retraction of the rear end of the cells induced by Yoda1 was proposed [55]. Another mechanism of Piezo-controlled cell motility was reported in Dictyostelium where Piezo channels were shown to control whether cells migrate with blebs or pseudopods [56]. The genetic depletion of Piezo1 expression can also have opposite effects on cellular motility: that is, the reduction of cell migration [57–59] or the increase of migratory properties of the cells after Piezo1 knockdown [60,61]. The variability of the effect of the manipulation of Piezo1 activity on cell migration is more likely explained by the coupling of Piezo1 with different intracellular signaling pathways in the cells.

Somatic stem cells permanently regulate their self-renewal and lineage commitment by combining different environmental signals to support tissue homeostasis. Although multiple chemical and biological signals that control stem cell behavior were recovered, whether stem cells can directly respond to mechanical stimuli remains unclear. This investigation has made some significant findings: first, Piezo1 was identified in eMSCs, and it acts as a molecular mechanism that contributes to mechano-induced Ca\(^{2+}\) entry in eMSCs. Second, ORAI1/STIM1/STIM2 proteins are expressed and mediate store-operated Ca\(^{2+}\) influx in eMSCs and act as a downstream mechanism that participates in Yoda1-induced Ca\(^{2+}\) entry. Third, both Piezo1-induced and store-operated Ca\(^{2+}\) entry suppress the eMSC migration, the rate of cell proliferation and induce cell cycle arrest without any effects on cell viability. We believe that the knowledge about the mechanisms underlying Piezo1-induced Ca\(^{2+}\) signaling and their role in eMSC migratory activity will help to develop novel strategies aimed at modulation of the regenerative potential of eMSC for tissue repairing.

4. Materials and Methods

4.1. Cell Cultures and Reagents

The endometrial mesenchymal stem cell (eMSC) line isolated from the desquamated endometrium of the menstrual blood of a healthy donor (line no. 2804, [11]) was a gift from the Department of Intracellular Signalling and Transport, Institute of Cytology, RAS. The cells had properties that are typical for MSCs, including fibroblast-like morphology, multipotency and the expression of standard surface markers [62]. Cells were grown in DMEM/F12 cultural medium (Gibco, Waltham, MA, USA) with 10% fetal bovine serum (HyClone, Logan, UT, USA) and 1% penicillin/streptomycin (Gibco, Waltham, MA, USA). The cells were maintained in culture flasks at 37 °C in 5% CO\(_2\). 2-[(2,6-dichlorophenyl)methylsulfanyl]-5-pyrazin-2-yl-1,3,4-thiadiazole (Yoda1) was purchased from Tocris (Tocris,
Abington, United Kingdom, cat. no. 5586). 2-aminoethyl diphenylborinate (2-APB) and thapsigargin (TG) were purchased from Sigma-Aldrich (Sigma-Aldrich, St.Louis, MO, USA, cat. no. D9754 and T9033, respectively).

4.2. Total RNA Extraction and Reverse Transcriptase (RT)-PCR

Total RNA was isolated using the RNeasy Mini Kit (Qiagen, Germantown, MD, USA). The PCR primers were designed using the GeneRunner v5.0.59 software. The primer sequences are represented in Table 1. To avoid false positive results due to genomic contamination of the samples, the primers spanned an intron at the genomic level. PCR was performed in the volume of 10 µL using 1 µL diluted (1:3) cDNA, 0.3 µM of each primer, 200 µM dNTPs, 2 mM MgCl₂, 1 unit Hot-Taq polymerase and 1 × Hot-Taq polymerase buffer (Sileks, Moscow, Russia). In negative control experiments, MMLV reverse transcriptase was omitted. PCR reaction products were subjected to electrophoresis on a 6% polyacrylamide gel and visualized by UV fluorescence after ethidium bromide staining.

Table 1. Primers for RT-PCR analysis.

| Gene   | Forward Primer | Reverse Primer |
|--------|----------------|----------------|
| hPIEZO1| 3′-CCAGAACAGGTATCGGAAG-5′ | 5′-TGCTGTACCACTGTACCTGCTG-3′ |
| hORAI1 | 5′-GACCTCGGCTCTGCTCTC-3′ | 5′-GGTGCTGACCATGGTTAC-3′ |
| hORAI2 | 5′-GCGGAAGCTCTACCTGAG-3′ | 5′-CCACATTGGGCAGGATGC-3′ |
| hORAI3 | 5′-CACGTCTGCCTTGCTCTC-3′ | 5′-ATGTTGCTCAGCCGTTCAATG-3′ |
| hSTIM1 | 5′-GGCAGTCCGTAACATCCAC-3′ | 5′-TTGTATACTTCTGATGACTTCC-3′ |
| hSTIM2 | 5′-ATGGTGGAATTGAAGTAGAGG-3′ | 5′-TTCCTTTGACATTGTTGTCTC-3′ |

4.3. Immunofluorescence

For immunofluorescent staining, the cells, preliminary plated on glass coverslips, 2–3 days before the experiments, were fixed with 3.7% paraformaldehyde in phosphate-buffered saline (1 × PBS, 10 min) and permeabilized with 0.25% Tween-20 (10 min at room temperature, RT). Non-specific binding of the antibodies (Abs) was blocked by incubation with 10% goat serum for 1 h at RT. Then cells were incubated with goat anti-rabbit polyclonal Abs against Piezo1 (Novus Bio, Centennial, CO, USA, cat. no. NBP1-78537) or rabbit polyclonal Abs against ORAI1 (Alomone Labs, Jerusalem, Israel, cat. no. ACC-060) or Abs against STIM1 (Alomone Labs, Jerusalem, Israel, cat. no. ACC-063) or STIM2 (Alomone Labs, Jerusalem, Israel, cat. no. ACC-064) at 4 °C overnight. Goat anti-rabbit (GAR)-Cy3 Abs (Santa Cruz, Dallas, TX, USA) were used for secondary detection. Cell nuclei were counterstained with DAPI (Sigma-Aldrich, St.Louis, MO, USA, final concentration was 0.1 µg/mL) and then the samples were mounted using Vectashield mounting medium (Vector Lab. Inc., Burlingame, CA, USA). As a control, staining of the cells only with fluorescent secondary Abs (GAR-Cy3, Santa Cruz, Dallas, TX, USA) was used. Images were obtained using the Olympus FV3000 laser scanning confocal inverted microscope (Olympus Corporation, Shinjuku, Tokyo, Japan) with 40×/1.3 NA oil immersion objective and a specific combination of laser/detectors.

4.4. Electrophysiology

Single currents were recorded using cell-attached configuration, as described previously [63]. Briefly, the patch-clamp set-up was based on the Axopatch 200B operational patch-clamp amplifier (Molecular Devices, San Jose, CA, USA) and Axon Digidata 1550A (Molecular Devices, San Jose, CA, USA) analog–digital converter controlled by a Windows-based personal computer with installed Axon PClamp 10.7 Software Suite (Molecular Devices LLC, San Jose, CA, USA) for data acquisition, processing and analysis. Patch pipettes were pulled from borosilicate glass capillaries (BF-150-86-10, Sutter Instruments, Novato, CA, USA) to a resistance of 10–15 MOhm when filled with the extracellular solution containing (in mM): 145 NaCl, 2 CaCl₂, 1 MgCl₂ and 10 HEPES/TrisOH. Potassium bath solution containing 145 KCl, 2 CaCl₂, 1 MgCl₂ and 10 HEPES/TrisOH was used to nullify
the resting membrane potential of the cells. The pH of all solutions was maintained at 7.3. All experiments were performed at RT. The single-channel recordings were processed and analyzed in Clampfit software (part of Axon PClamp 10.7 Software Suite). Single-channel conductance values were defined by the slope of the current–voltage relationship (I–V) after linear approximation. Averaged conductance values are given as the mean ± SEM (n—number of experiments).

4.5. Ca\textsuperscript{2+} Imaging

eMSCs were seeded on cover slides 2–3 days before Ca\textsuperscript{2+} measurements. On the day of the experiment, cells were washed with a serum-free medium and loaded for 35 min with a 4 \textmu M Fura-2AM probe (Thermo Fisher Scientific, Waltham, MA, USA) in the dark at RT. Then, cells were washed, and the cover slides were transferred into a perfusion chamber. Cell imaging was obtained using an AxioObserverZ1 inverted microscope (Carl Zeiss MicroImaging GmbH, Oberkochen, Germany) with a Carl Zeiss Fluar 40×/1.3 Ph3 NA oil objective. Fura-2AM fluorescence was excited every 5 s sequentially by the light of 340 nm or 380 nm from an illuminator with a Lambda DG-4 high-speed wavelength switcher (Sutter Instrument Co., Novato, CA, USA). Filter set 21 HE (Carl Zeiss MicroImaging GmbH, Oberkochen, Germany) was used (excitation BP 340/30, emission BP 510/90). “Physiology” plugin of AxioVision 4.8.2 software (Carl Zeiss MicroImaging GmbH, Oberkochen, Germany) was used for data acquisition and analysis. Calcium imaging recordings were subjected to the subtraction of the background fluorescence. Fura-2AM 340/380 nm ratio values are given as mean ± SD (n—number of the cells).

4.6. Quantification of Cell Migration (Wound Healing Assay)

For the wound healing assay, trypsinized eMSCs were seeded into Ibidi Culture-Inserts 2 Well (Ibidi GmbH, Gräfelfing, Germany, total number of cells was 70,000 per insert) pre-installed in 24-well culture plates to form the experimental wounds (gaps of the standardized sizes). The cells were cultured for a further 24 h to allow cell spreading, then, the inserts were removed, and the plates were transferred to the stage of a Carl Zeiss AxioObserver Z1 (Carl Zeiss Microimaging GmbH, Oberkochen, Germany) microscope equipped with the set-up for long-term live-cell imaging under stable culture conditions (controlled humidity, 37 °C and 5% CO\textsubscript{2}). The images of the experimental wounds were acquired automatically (every 2 h; total time of the experiment was 48 h) using the “Multidimensional Acquisition” plugin of Carl Zeiss Axiovision 4.8.2 Software (Carl Zeiss MicroImaging GmbH, Oberkochen, Germany). The sizes of experimental wounds were calculated manually using the “Measure Area” tool in ImageJ software (NIH, Bethesda, MD, USA), and the wound sizes were normalized to the size of the initial wound (at 0 h starting measurement). Wound sizes are shown as mean ± SEM (n—number of independent experiments). Data were compared using a two-sample Student’s t-test (for data with unequal variances, Welch’s correction was introduced); p < 0.05 was considered as significant.

4.7. FACS Analysis

Detached with trypsin/EDTA solution, eMSCs were suspended in growth medium and utilized for viability and cell cycle analysis as described previously [64]. Briefly, 0.05 mg/mL of propidium iodide (PI) was added to the eMSCs and exposed to flow cytometry (FACS) analysis. The cells gated as PI-negative were used for growth curve generation. For cell cycle analysis, saponin (0.2 mg/mL), RNAse (0.25 mg/mL) and PI (0.05 mg/mL) were applied to cell suspension, mixed gently and incubated for 1 h in the dark at RT. No less than 3000 events were gathered for viability assay and 15,000 events for cell cycle analysis. Cytometric analysis was conducted with CytoFLEX S flow cytometer (Beckman Coulter, Brea, CA, USA) equipped with Cytexpert software (version 2.0). Data were compared using a two-sample Student’s t-test (p < 0.05 was considered as significant) and are presented as mean ± SD.
4.8. Statistics

Statistical analysis was performed using GraphPad Prism 6.0 software (GraphPad Software, San Diego, CA, USA). All data were tested for normal distribution using Shapiro–Wilk’s test and homogeneity of variances (Levene’s test). The particular, statistical criteria used for means comparison are described in the relevant sections of Materials and Methods; \( p < 0.05 \) was considered as significant.

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