# Effects of human umbilical cord mesenchymal stem cells derived from exosomes on migration ability of endometrial glandular epithelial cells

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Abstract. The present study aimed to investigate the effects of human umbilical cord mesenchymal stem cells (Huc-MSCs)-derived exosomes on the migratory abilities of endometrial glandular epithelial cells, and to evaluate the underlying mechanism from the perspective of epithelial-mesenchymal transition (EMT). Huc-MSCs were prepared from human umbilical cord, and eutopic endometrial glandular epithelial cells were isolated from patients with endometriosis. The exosomes derived from Huc-MSCs (Huc-MSCs-exo) were prepared using an exosome extraction kit. The endometrial glandular epithelial cells were randomly divided into two groups: Huc-MSCs-exo and control. Cell migratory ability was assessed and western blotting was used to detect the expression levels of EMT. The results of the present study demonstrated that Huc-MSCs-exo treatment significantly enhanced the migration of endometrial glandular epithelial cells from patients with endometriosis (P<0.05). The present study also demonstrated that treatment with Huc-MSCs-exo inhibited the expression levels of E-cadherin and promoted the expression levels of Vimentin and N-cadherin at both the mRNA and protein level. The results of the current study indicate that Huc-MSCs-exo enhance the migratory ability of endometrial glandular epithelial cells via promotion of EMT.

# Introduction

Although endometriosis is considered to be a benign disease, its characteristics (including ectopic invasion and high recurrence) endow it with the biological behavior of malignant tumors (1,2), of which ovaries and uterosacral ligaments are the most common locations. Symptoms of endometriosis include chronic pelvic pain, infertility and adnexal cystic mass (3,4). Adhesion-invasion-implantation is the pathophysiological process underlying endometriosis and involves early cell detachment from the primary lesion and dissemination. At the later stages, circulating cells anchor and proliferate in distant tissues to form heterotopic lesions (5).

Mesenchymal stem cells (MSCs) are located in a number of tissues, such as adipose tissue and bone marrow (6). MSCs can migrate from the original tissue to the site of pathophysiological changes in response to inducing factors, such as intercellular adhesion molecule-1 (7). Human umbilical cord MSCs (Huc-MSCs) are active throughout pregnancy. Exosomes are membrane bound extracellular vesicles that are produced in the endosomal compartment of eukaryotic cells, they serve as important biomarkers to identify different diseases and as therapeutic targets for diseases (8). Exosomes derived from MSCs contain abundant biological information, including proteins, such as membrane receptors, ribosomes and genetic information. These regulate tissue regeneration and repair, immunoregulation, cell growth and differentiation regulation (9-11). To the best of our knowledge, however, the effects of Huc-MSCs on the biological activities of endometrial glandular epithelial cells derived from patients with endometriosis have not previously been reported.

Epithelial-mesenchymal transition (EMT) refers to morphological and phenotypic transformation of epithelial cells to mesenchymal cells in response to stimulation of certain physiological or pathological changes, such as the loss of cell polarity, decreased contact with surrounding cells and the extracellular matrix, and increased migration and mobility (12). EMT is involved in early embryonic development and organogenesis, as well as wound healing (13). In addition, the metastasis of numerous types of malignant tumors, including lung cancer, is associated with EMT (14). EMT is a complex process by which epithelial cells acquire the characteristics of invasive mesenchymal cells (15).

The present study aimed to investigate whether exosomes derived from MSCs affect the migratory ability of endometrial glandular epithelial cells. In order to identify the underlying mechanism, the present study evaluated the effects of exosomes on EMT in endometrial glandular epithelial cells.

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#### Materials and methods

*Tissue samples*. The *in situ* endometrial tissue from five patients with endometriosis (age range, 31-42 years) and human umbilical cord tissue from six normal delivery woman (age range, 25-32 years) were provided by The Second Affiliated Hospital of Nanchang University (Nanchang, China) between June 2018 and July 2018. The experimental protocols were approved by the Ethics Committee of Nanchang University (China). All patients agreed to the use of their samples in scientific research and written informed consent was obtained from all patients.

Preparation of endometrial glandular epithelial cells. Ectopic endometrial tissue from patients with endometriosis was collected and cleaned using D-Hank's solution (cat. no. H1045; Beijing Solarbio Science & Technology Co., Ltd.) containing antibiotics, and blood vessels and impurities in the tissue were shaved using a scalpel. Ophthalmic scissors were used to cut tissues into 1 mm<sup>3</sup> blocks in a sterile Petri dish and the samples were transferred to a sterile centrifugal tube. Collagenase IV (0.1%; Sigma-Aldrich; Merck KGaA) was added to the centrifugal tube and incubated at 37°C for 40-60 min. The digestion was terminated by adding DMEM with 20% FBS (Hyclone; Cytiva) after intermittent oscillation. The solution was filtered through 100- and 400-mesh screens to remove large tissue fragments. The large tissue fragments on the surface of 100-mesh screen were transferred into a centrifugal tube, and collagenase (0.1%) was added for a secondary digestion at 37°C for 5 min. Cell suspension was collected, centrifuged (1,000 x g at room temperature for 10 min), and suspended cells were inoculated on the culture plate. The cells were incubated at 37°C with 5% CO<sub>2</sub>.

Preparation of Huc-MSCs-derived exosomes (Huc-MSCs-exo). The blood vessels in the Huc tissue were removed using a scalpel under a stereomicroscope, and tissues were washed with PBS, cut into 1 mm<sup>3</sup> blocks and digested with diluted trypsin (PBS mixed with trypsin; 1:1) overnight at 4°C. Tissue in the Petri dish was transferred to a 50 ml Eppendorf tube and placed in a 37°C water bath for 15 min. DMEM with 20% FBS was added to the Petri dish and cultured in an incubator for 15 min at 37°C and 5% CO<sub>2</sub>. The cells were incubated with the following antibodies: Isothiocyanate (FITC) anti-CD34 (1:100; cat. no. 343604; BioLegend, Inc.), FITC anti-CD44 and FITC anti-CD45 (1:100; cat. nos. ab46793 and ab134199, respectively; both purchased from Abcam) at room temperature in the dark for 10 min and detected using a NovoCyte™ flow cytometer (NovoCyte 2060R; ACEA Bioscience, Inc; Agilent).

*Extraction of Huc-MSCs-exo.* Following 48 h starvation with FBS-free medium at 37°C, exosome extraction kit was used to extract exosomes (cat. no. E1310; Bioruo) according to the manufacturer's instructions. Cells and debris were removed by centrifugation  $(2,000 \text{ x g}; 4^{\circ}\text{C}; 10 \text{ min})$ . The isolated exosomes were observed using transmission electron microscopy. The exosomes were fixed using 2.5% glutaraldehyde, phosphoric acid buffer preparation for 2 h at room temperature. After embedding, sections were cut (thickness, 70 nm) and stained

with 3% uranium acetate and lead citrate for 10 min at room temperature. The slides were observed using transmission electron microscopy [JEM-1230 (80KV); JEOL, Ltd.], at magnification x1,000. A total of 250  $\mu$ g Huc-MSCs-exo was labeled using PKH Lipophilic Membrane Dyes (cat. no. PKH67GL; Sigma-Aldrich, Merck KGaA) according to the manufacturer's instructions. PKH67-labeled Huc-MSCs-exo were centrifuged (40,000 x g; 4°C; 70 min) and suspended in PBS (50  $\mu$ l). Then, 2  $\mu$ l PKH67-labeled Huc-MSCs-exo solution was added to cells, which were incubated at 37°C for 0.5, 2.0 and 4.0 h. The cells were imaged using Zeiss confocal laser scanning microscopy (LSM710; Zeiss GmbH), at magnification x200.

*Experimental groups*. The samples were divided into two groups (n=6): Control and Huc-MSCs-exo treatment ( $10 \mu g/ml$ ) group. Subsequent experiments were performed 24 h after treatment.

*Transwell assay.* Following treatment with Huc-MSCs-exo for 24 and 48 h, 3x10<sup>3</sup> endometrial glandular epithelial cells were seeded in the upper chamber of Transwell plates (HyClone; GE Healthcare Life Sciences) with serum-free DMEM. The lower chamber contained DMEM with 10% FBS (Hyclone; Cytiva). One day after seeding, the cells in the lower chamber were fixed with 4% paraformaldehyde for 30 min at room temperature and stained with 1% crystal violet (Beijing Solarbio Science & Technology Co., Ltd.) for 5 min at room temperature. Images were captured using a light microscope (magnification, x200).

*Wound healing assay.* Endometrial glandular epithelial cells were seeded in 6-well plates (3x10<sup>3</sup> cells/well) in DMEM with 20% FBS, and treated with Huc-MSCs-exo for 24 and 48 h. The cell monolayer was scratched. After 24 and 48 h incubation at 37°C, images were captured using a light microscope (Olympus Corporation), at magnification x100.

Reverse transcription-quantitative (RT-q)PCR. At 24 h post-treatment, total RNA was extracted from endometrial glandular epithelial cells using an Ultrapure RNA extraction kit (CoWin Biosciences), according to the manufacturer's protocols. The purity of RNA was assessed by measuring optical density at 280/260 nm using a spectrometer (LASPEC). RNA (1  $\mu$ g) was reverse transcribed into cDNA using an Avian Myeloblastosis Virus Reverse-Transcriptase kit (cat. no. KL041; Shanghai Kang Lang Biological Technology Co., Ltd.). The reaction system included 9.5  $\mu$ l RNase-free dH<sub>2</sub>O, 1.0 µl cDNA/DNA, 2.0 µl primers and 12.5 µl UltraSYBR mixture (cat. no. 00081405; CoWin Biosciences) and PCR was performed using the following thermocycling conditions: Initial denaturation at 95°C 10 min; followed by 40 cycles of denaturation at 95°C for 10 sec, annealing at 58°C for 30 sec and extension at 72°C for 30 sec; and final extension at 72°C for 3 min. GAPDH was used as a reference gene. The following primers were used: E-cadherin forward, 5'-CTCACATTTCCCAACTCCTCT-3' and reverse, 5'-TGT CACCTTCAGCCATCCT-3'; Vimentin forward, 5'-GGA TTCACTCCCTCTGGTTG-3' and reverse, 5'-TGATGCTGA GAAGTTTCGTTG-3'; N-cadherin forward, 5'-GCTTAT



Figure 1. Labeling of Huc-MSCs-exo in endometrial glandular epithelial cells. (A) Flow cytometry for Huc-MSCs. (B) Immunohistochemistry for endometrial glandular epithelial cells. Scale bar, 100  $\mu$ m. (C) Morphology of Huc-MSCs-exo detected using transmission electron microscopy. Scale bar, 50 nm. (D) Labeling of HucMSCs-exo in endometrial glandular epithelial cells. Scale bar, 100  $\mu$ m. Huc-MSCs-exo, human umbilical cord mesenchymal stem cell-derived exosomes.

CCTTGTGCTGATGTTT-3' and reverse, 5'-GTCTTCTTC TCCTCCACCTTCT-3'; and GAPDH forward, 5'-CAATGA CCCCTTCATTGACC-3' and reverse, 5'-GAGAAGCTT CCCGTTCTCAG-3'.

Western blotting. At 24 h post-treatment with Huc-MSCs-exo, protein was extracted from cells using the triplePrep kit (cat. no. 28-9425-44; ReadyPrep; GE Healthcare Life Sciences). The protein levels were quantified using a bicinchoninic acid protein assay kit (Beyotime Institute of Biotechnology), according to the manufacturer's instructions. A total of 25  $\mu$ g/lane protein was separated via SDS-PAGE on a 10% gel and transferred onto nitrocellulose membranes. The membranes were blocked using 5% skimmed milk at room temperature for 2 h and incubated overnight at 4°C with the following primary antibodies: Rabbit polyclonal anti-E-cadherin (1:1,500; cat. no. AF0131; Affinity Biosciences), anti-GAPDH (1:1,000; cat. no. AG019; Beyotime Institute of Biotechnology), anti-Vimentin and anti-N-cadherin (1:3,000 and 1:1,000, respectively; cat. nos. ab92547 and ab76057, respectively; both Abcam). After washing with 0.1 M PBS, the membranes were incubated with the secondary antibody (HRP-labeled goat anti-rabbit IgG; cat. no. A16104; Thermo Fisher Scientific, Inc.) at 4°C for 2 h. The blots were visualized using an electrochemiluminescence kit (Thermo Fisher Scientific, Inc.). The densities of the blots were quantified using the Quantity One software (version 4.62; Bio-Rad Laboratories, Inc.).

Immunohistochemistry. The cells were fixed in paraformaldehyde for 30 min at room temperature. Staining was performed using monoclonal antibodies against N-cadherin, Vimentin (both 1:100; cat nos. ab76057 and ab92547, respectively; both Abcam) and E-cadherin (1:200; cat. no. AF0131; Affinity Biosciences). Endogenous peroxidase activity was blocked with 3% (v/v)  $H_2O_2$  for 5 min at room temperature. The slides were incubated with the aforementioned primary antibodies overnight at 4°C, followed by incubation with horseradish peroxidase-labeled goat anti-rabbit IgG secondary antibody (1:10,000; cat. no. A16104SAMPLE; Thermo Fisher Scientific, Inc.) for 30 min at room temperature. Cells were stained with 3,3'-diaminobenzidine chromogen for 3 min at room temperature. Images were captured using a light microscope (BX51; Olympus Corporation), at magnification, x200.

Statistical analysis. Data are expressed as the mean  $\pm$  standard deviation of six independent experiments. All statistical analysis was performed using unpaired Student's t-test using GraphPad Prism (version 7.0; GraphPad Software, Inc.). P<0.05 was considered to indicate a statistically significant difference.



Figure 2. Huc-MSCs-exo facilitate wound healing ability of endometrial glandular epithelial cells. Left, representative images of cell migration (magnification, x100); right, quantification data. \*P<0.05 vs. control. Huc-MSCs-exo, human umbilical cord mesenchymal stem cell-derived exosomes.



Figure 3. Huc-MSCs-exo facilitate migratory ability of endometrial glandular epithelial cells. Arrows indicate migratory cells. \*P<0.05 vs. control. Huc-MSCs-exo, human umbilical cord mesenchymal stem cell-derived exosomes.

# Results

*Identification of Huc-MSCs and endometrial glandular epithelial cells.* CD44, CD34 and CD45 expression levels in the isolated cells were 92.40, 6.80 and 0.27%, respectively

(Fig. 1A). These data indicated that the isolated cells were CD44<sup>+</sup>, CD34<sup>-</sup> and CD45<sup>-</sup>, which revealed the successful isolation of umbilical cord-derived MSCs. Cells expressed vimentin (dark brown), which indicated that endometrial glandular epithelial cells were successfully isolated (Fig. 1B).



Figure 4. Huc-MSCs-exo increase expression levels of N-cadherin and Vimentin, and decrease E-cadherin expression levels. (A) N-cadherin, Vimentin and E-cadherin expression levels detected by immunohistochemistry. Scale bar, 100  $\mu$ m. (B) mRNA expression levels detected via reverse transcription-quantitative PCR. (C) Protein expression levels detected via western blotting. \*P<0.05 vs. control. Huc-MSCs-exo, human umbilical cord mesenchymal stem cell-derived exosomes.

Labeling of Huc-MSCs-exo in endometrial glandular epithelial cells via transmission electron microscopy. Exosomes exhibited typical round or oval cup-shaped structures with complete morphology (Fig. 1C). Huc-MSCs-exo entered endometrial glandular epithelial cells over time (Fig. 1D). At 4 h, the labeled exosomes were notably expressed in the cytoplasm.

Huc-MSCs-exo facilitate the wound healing ability of endometrial glandular epithelial cells. As shown in Fig. 2, the migratory ability of endometrial glandular epithelial cells treated with Huc-MSCs-exo was significantly enhanced at 24 and 48 h compared with that of the control group (P<0.05), which demonstrated that Huc-MSCs exosomes promote the migration of endometrial glandular epithelial cells. *Huc-MSCs-exo facilitate the migratory ability of endometrial glandular epithelial cells.* Compared with control group, the migratory ability of HucMSCs-exo-treated cells significantly increased at 24 and 48 h (P<0.05; Fig. 3), which demonstrated that Huc-MSCs-exo promote the migratory ability of endometrial glandular epithelial cells.

Huc-MSCs-exo increase expression levels of N-cadherin and Vimentin, and decrease E-cadherin expression levels. Immunohistochemical staining demonstrated that the relative expression levels of N-cadherin and Vimentin increased and the expression level of E-cadherin decreased in the Huc-MSCs-exo group compared with the control group (P<0.05; Fig. 4A). RT-qPCR also demonstrated that Huc-MSCs-exo treatment promoted expression levels of N-cadherin and Vimentin and decreased expression levels of E-cadherin at the mRNA level (Fig. 4B). This was consistent with the results of the western blotting analysis (Fig. 4C).

# Discussion

The present study demonstrated that Huc-MSCs-exo promoted the migration of endometrial glandular epithelial cells from patients with endometriosis at both 24 and 48 h. It was also demonstrated that treatment with Huc-MSCs-exo inhibited expression levels of E-cadherin and promoted expression levels of Vimentin and N-cadherin. The results of the present study indicate the potential function of Huc-MSCs-exo in patients with endometriosis.

Endometriosis is a common gynecological disease, primarily occurring in women of child-bearing age (25-45 years). It is estimated that 10.8 million individuals were affected globally as of 2015 (16). The primary clinical symptoms include dysmenorrhea, increased menstrual volume and infertility, which may adversely affect the quality of life of patients (17). In the present study, endometrial glandular epithelial cells were isolated and identified via immunohistochemistry. MSCs are adult stem cells that originate from the embryonic mesoderm and possess strong self-renewal ability and multi-directional differentiation potential (18). Numerous studies have demonstrated that MSCs can serve as the host of primary or metastatic tumors and promote the formation of tumor microenvironment (19,20). Studies have demonstrated that MSCs can either promote or inhibit the growth of tumors (21,22). MSCs may promote the development of cancer via EMT, which is characterized by the transformation of cells from epithelial to mesenchymal phenotype (23). In the present study, Huc-MSCs were isolated from Huc tissues and identified via flow cytometry using specific antibodies as previously described (23).

Exosomes from numerous cell sources contain biologically active components, such as cell membrane molecules and cytoplasmic proteins (24). Intercellular information is transferred via fusion with the cell membranes of adjacent cells (25). Herrera et al (26) demonstrated that exosomes derived from human stem cells accelerate liver regeneration in hepatectomized rats. Exosomes are considered to be potential mediators for inducing peripheral tolerance and regulating the immune response (27). Exosomes are considered to be mediators of intercellular communication, and influence target receptor cells by inducing intracellular signal transduction or endowing donor cells with novel substances, such as protein or mRNA (9-11). In the present study, exosomes were identified using transmission electron microscopy. It was demonstrated that the exosomes were round or elliptical in shape and conformed to typical morphological characteristics. The labeling of endometrial glandular epithelial cells by exosomes demonstrated that the majority of the exosomes entered cells, allowing them to exert their function.

Endometriosis is a non-cancerous disease with invasive ability. The invasiveness of endometrial cells serves a key role in the occurrence and development of endometriosis. Huc-MSCs-exo promote the migratory and invasive abilities of A549 cells (23). Numerous studies have demonstrated that MSC-exo exerts both anti-tumor and pro-tumor effects in human breast, ovarian, gastric and nasopharyngeal cancer, as well as in multiple myeloma, osteosarcoma and rat liver cancer (28-36). MSC-exo also promotes migration and invasion of human lung cancer cells (23). The results of the present study demonstrated that Huc-MSCs-exo promote the migration of endometrial glandular epithelial cells.

EMT refers to the loss of epithelial characteristics and acquisition of mesenchymal phenotypes. When cells undergo EMT, they detach from other cells, decrease expression levels of epithelial markers and acquire mesenchymal characteristics, such as enhanced mobility and invasiveness (37,38). Zhu et al demonstrated that human MSC-conditioned medium induces EMT in cancer cells (39,40). The active component of conditioned medium is exosome from mesenchymal stem cells (39,40). A previous study demonstrated that expression levels of EMT markers are altered by exosomes in vivo (36). Bartley et al (41) demonstrated that expression levels of E-cadherin are absent in certain glandular epithelial cells in ectopic lesions of patients with endometriosis, which indicates that EMT may also play a key role in the formation and progression of endometriosis. EMT-associated transcription factors can directly or indirectly inhibit expression levels of E-cadherin and promote expression levels of N-cadherin and Vimentin in mesenchymal phenotypes, thereby promoting EMT (42). The results of the present study demonstrated that Huc-MSCs-exo promote EMT and upregulate expression levels of N-cadherin and Vimentin, while inhibiting expression levels of E-cadherin. This is consistent with the aforementioned results and supports the hypothesis that Huc-MSCs-exo alter expression levels of EMT-associated genes, thus enhancing the migratory and invasive abilities of endometrial glandular epithelial cells.

In the present study, Huc-MSCs-exo promoted the migratory ability of endometrial glandular epithelial cells from patients with endometriosis. The current results indicate the potential application of Huc-MSCs-exo in the treatment of endometriosis. Further research is required to quantify the dose effect of Huc-MSCs-exo and to investigate the underlying mechanism. Huc-MSCs-exo function should also be further evaluated in an *in vivo* model.

In conclusion, Huc-MSCs-exo promote the migratory ability of endometrial glandular epithelial cells, potentially via promoting EMT in endometrial glandular epithelial cells.

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# Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

#### Authors' contributions

YF, FZ and YZ performed the experiments and analyzed the data. YF and BT designed the study and wrote the manuscript. All authors read and approved the final manuscript.

# Ethics approval and consent to participate

The experimental protocols were approved by the Ethics Committee of Nanchang University (approval no. 2018071401).

#### Patient consent for publication

All patients agreed to the use of their samples in scientific research and written informed consent was obtained from all patients.

#### **Competing interests**

The authors declare that they have no competing interests.

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