# L1CAM promotes epithelial to mesenchymal transition and formation of cancer initiating cells in human endometrial cancer

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Abstract. Identification of novel factors critical for epithelial to mesenchymal transition (EMT) and cancer initiating cell (CIC) formation may aid in the identification of novel therapeutics for the treatment of endometrial cancer. The present study demonstrated that L1 cell adhesion molecule (CAM) is critical for EMT and formation of CICs in endometrial cancer. Overexpression of L1CAM may promote EMT with increased formation of CICs in HEC-1A endometrial cancer cells. CICs and mesenchymal status resist chemotherapeutic drugs and may regenerate the various cell types in tumors, thereby resulting in relapse of the disease. The present study demonstrated that overexpressing L1CAM promoted paclitaxel resistance and regulated paclitaxel resistance-associated microRNA expression in HEC-1A cells. Furthermore, it was demonstrated that overexpressing L1CAM promoted anoikis resistance in HEC-1A cells. This link between L1CAM and EMT/CICs may provide a novel target for advancing anticancer therapy.

## Introduction

Endometrial carcinoma (EC) is the most common gynecologic malignancy and is associated with a poor prognosis when diagnosed at an advanced stage (1). Endometrial cancer is traditionally classified into type I and type II subtypes (2). Type I cancers account for 80-85% of EC cases, are of endometrioid histology, more often well differentiated and associate with favorable prognosis (2). In contrast the type II

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*Abbreviations:* CICs, cancer initiating cells; EMT, epithelial to mesenchymal transition; EC, endometrial cancer

*Key words:* endometrial cancer, cancer initiating cells (CICs), epithelial to mesenchymal transition (EMT), L1CAM, paclitaxel resistance

cancers are non-endometrioid carcinomas, poorly differentiated and associate with poorer survival (2). However, patients with deep myometrial invasion, poor differentiation, serous or clear cell histology or extension of disease to other organs or lymph nodes within the pelvic region are at higher risk for disease recurrence (3,4). Therefore, it is imperative to find new therapeutic targets to elaborate the molecular mechanisms underlying progression of endometrial carcinogenesis.

L1 cell adhesion molecule (L1CAM, CD171) is a 200-220-kDa transmembrane glycoprotein composed of 6 immunoglobulin-like domains, 5 fibronectin-type III domains, a transmembrane stretch, and a short cytoplasmic tail (5). L1CAM was originally identified as a neural cell adhesion molecule in the central nervous system that plays an important role in initiating cerebellar cell migration and neurite outgrowth (6). L1CAM expression is also found in other cell types such as lymphoid and myelomonocytic cells, kidney tubule epithelial cells, and intestinal crypt cells (7-10). In addition, L1CAM expression has been identified in a variety of tumor types and correlates with poor prognosis and metastasis (11). L1CAM functions mostly in proliferation, migration, invasion, and survival through L1CAM homophilic interaction or heterophilic interactions with other cell adhesion molecules, integrins, or growth factor receptor, while the cellular properties are not homogeneous among different types of cancers (12). Recently, it has been reported that L1CAM was involved in progression of endometrial cancer (13).

Cyclophilin A (CypA) is a highly abundant protein, accounting for up to ~0.6% of the total cytosolic protein content (14). CypA is involved in a growing number of biological processes, including protein folding, signal transduction, viral infection, trafficking, receptor assembly, immune response, and transcription regulation (15). Although several proteins have been identified to interact with CypA (16-19), the underlying mechanism of the CypA action and the physiological implications of the interactions remain in most cases unknown. CypA exhibits peptidyl-prolyl *cis-trans* isomerase (PPIase) activity by catalyzing *cis-trans* isomerization of peptide bonds preceding proline residues (20). CypA can in principle act as an enzyme or a binding partner (21) in mediating the biological processes.

In this study, we showed that L1CAM promotes EMT with increased characteristics of CICs and paclitaxel resistance in human endometrial cancer.

## Materials and methods

*HEC-1A cells line.* HEC-1A cells were obtained from Peking Union Medical College (Beijing, China). Briefly, cells were maintained in Dulbecco's modified Eagle's medium (DMEM; Sigma, Shanghai, China) containing 10% fetal bovine serum (FBS; Shanghai ExCell Biology, China) and 100 mg/ml penicillin and streptomycin (Gibco, Shanghai, China) at 37°C in a humidified atmosphere with 5% CO<sub>2</sub>.

LICAM expressing plasmids/empty vectors and transfection experiments. LICAM expressing plasmids and empty vectors (pcDNA3.1) were obtained from Tiangen (Beijing, China). Transfections were performed with Lipofctamine 2000 transfection reagent (Invitrogen, Carlsbad, USA) following the manufacturers' protocols.

Western blot analysis. It was performed as described previously (22,23). Total protein was prepared using extraction buffer comprising NaCl/P<sub>i</sub> containing 0.5% Triton X-100, 1 mM EDTA, 1 mM phenylmethyl sulfonyl fluoride, and complete protease inhibitors (Roche). The concentration of each protein lysate was determined using a BCA<sup>™</sup> protein assay kit (Thermo Scientific, Rockford, IL, USA). Equal amounts of total protein were subjected to 12% SDS/PAGE. Then samples were transferred to nitrocellulose membranes and blocked for 60 min at room temperature in 5% skim milk powder (w/v) in NaCl/P<sub>i</sub>. The membranes were immunoblotted using primary anti-body anti-L1CAM (1:500; Abcam, Cambridge, MA, USA), anti-E-cadherin (1:500; Abcam, Cambridge, MA, USA), anti-Vimentin (1:500; Abcam, Cambridge, MA, USA), anti-Musashi-1 (1:500; Abcam, Cambridge, MA, USA), anti-CD133 (1:500; Abcam, Cambridge, MA, USA), anti-Cyclophilin A (1:500; Abcam, Cambridge, MA, USA) and anti-β-actin (1:500; Abcam, Cambridge, MA, USA) overnight at 4°C, anti-rabbit secondary antibodies (1:10,000; Abcam, Cambridge, MA, USA) were used for 30 min at room temperature. The specific proteins were visualized by Odyssey<sup>TM</sup> Infrared Imaging System (Gene Company, Lincoln, NE, USA). β-actin expression was used as an internal control to show equal loading of the protein samples.

Immunofluorescence staining. It was performed as described previously (24,25). Cells were plated on glass coverslips in six-well plates and transfected as indicated. At 48 h after transfection, the cells were fixed in 4% paraformaldehyde for 15 min, and then blocked with goat serum blocking solution for 20 min at room temperature. Coverslips were stained with the mentioned antibody mentioned anti-L1CAM antibodies (1:500; Abcam, Cambridge, MA, USA), After washing three times with NaCl/P<sub>i</sub>, cells were incubated with appropriate secondary antibodies (Abcam, Cambridge, MA, USA) for 30 min at 37°C. 4'6-diamidino-2-phenylindole (DAPI) staining (blue) was used to indicate nuclei. Microscopic analysis was performed with a confocal laser-scanning microscope (Leica Microsystems, Bensheim, Germany). Fluorescence intensities were calculated from a few viewing areas for 300 cells per coverslip and analyzed by ImageJ 1.37v software (http://rsb.info.nih.gov/ij/index.html).

*Quantitative real-time RT-PCR (qRT-PCR).* Quantitative real-time RT-PCR were described before (21). The specific

primer sets for PCR were as follows: GAPDH, forward primer: 5'-GAAGGTGAAGGTCGGAGTCA-3', and reverse primer 5'-GAAGATGGTGATGGGATTTC-3'; E-Cadherin, forward primer 5'-TCAACGATCCTGACCAGCAGTTCG-3' and reverse primer 5'-GGTGAACCATCATCTGTGGGCGATG-3'; N-cadherin, forward primer 5'-CATCCCTCCAATCAACTT GC-3' and reverse primer 5'-ATGTGCCCTCAAATGAAA CC-3'; Vimentin, forward primer 5'-GACAATGCGTCTCTG GCACGTCTT-3' and reverse primer 5'-TCCTCCGCCTCC TGCAGGTTCTT-3'; ZEB1, forward primer 5'-TAGGTGCC CCCTGTGCAGTT-3' and reverse primer 5'-TAGGAGCCA GAATGGGAAAAG-3'. GAPDH was a loading control.

MTT assay. The proliferation of cells was assessed by the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium (MTT) assay (Sigma, St Louis, MO, USA). The MTT analysis was performed as described previously (26-29). In brief, the cells were plated in 96-well plates in Dulbecco's modified Eagle's medium containing 10% fetal bovine serum at a density of 8x10<sup>3</sup> cells per well at 37°C in a 5% CO<sub>2</sub> incubator for 12 h. Cells were transfected with L1CAM expressing plasmid or empty vectors for 24 h and then were treated with different doses of paclitaxel (10<sup>-4</sup>-10<sup>2</sup>). After 24 h, MTT (5 mg·ml<sup>-1</sup>) was added to the wells (20  $\mu$ l per well). The plates were incubated in a cell incubator for 4 h, then the supernatant was removed and  $150 \,\mu l$ of dimethyl sulfoxide was added to each well. After incubation for 10 min, the absorbance of each well was measured using a Synergy<sup>™</sup> 4 (BioTek Instruments, Winooski, VT, USA) with a wavelength of 570 nm, with the reference wavelength set at 630 nm. Absorbance was directly proportional to the number of survival cells.

*Sphere formation assay.* It was performed as described previously (30). Cells (10<sup>3</sup>/ml) in serum-free RPMI1640/1 mM Na-pyruvate were seeded on 0.5% agar precoated 6-well plates. After 1 week, half the medium was exchanged every third day. Single spheres were picked and counted.

Anoikis assays. It was performed as described previously (31). Anoikis resistance was evaluated by seeding 7.5x10<sup>4</sup> cells in ultralow attachment plates (Corning). After 24 h of anchorage-independent culture, cells were transfected as indicated and resuspended in 0.4% trypan blue (Sigma, St. Louis, MO, USA) and cell viability was assessed.

*miRNA microarray*. It was performed as described previously (32). Total RNA from cultured cells, with efficient recovery of small RNAs, was isolated using the mirVana miRNA Isolation Kit (Ambion, Austin, TX, USA). cRNA for each sample was synthesized by using 3' IVT EXPRESS KIT (Affymetrix, Santa Clara, CA, USA) according to the manufacturer's protocols. The purified cRNA was fragmented by incubation in fragmentation buffer (provided in the 3'IVT express kit) at 95°C for 35 min and chilled on ice. The fragmented labeled cRNA was applied to MicroRNA2.0 Array (Affymetrix, Santa Clara, CA, USA) and hybridized in Genechip hybridization oven 640 (Affymetrix, Santa Clara, CA, USA) at 45°C for 18 h. After washing and staining in Genechip fluidics station 450 (Affymetrix, Santa Clara, CA, USA), the arrays were scanned by using Genechip scanner 3000 (Affymetrix, Santa Clara, CA, USA). The gene expressions levels of samples were normalized and compared by using Partek GS 6.5 (Partek, Inc, St. Louis, MO, USA). Average-linkage hierarchical clustering of the data was applied by using the Cluster [Eisen *et al* (33), Stanford, Stanford University, CA, USA; http://rana.lbl.gov] and the results were displayed by using TreeView [Eisen *et al* (33), Stanford, Stanford University, CA, USA; http://rana.lbl.gov].

Statistical analysis. Results were analyzed using SAS software (9.4). Data were presented as mean  $\pm$  standard error of the mean (SEM) of separate experiments (n=3). P-values less than 0.05 were considered to be significant.

# Results

LICAM promotes EMT in endometrial cancer HEC-1A cells. To investigate whether L1CAM can affect epithelial or mesenchymal status of HEC-1A cells, we performed western blot to test whether L1CAM expressing plasmids could express L1CAM protein in HEC-1A cells. The results of western blot showed that L1CAM expressing plasmids can significantly up-regulate L1CAM protein expression in the cells (Fig. 1A). To determine whether L1CAM can promote EMT, we transfected HEC-1A cells with L1CAM expressing plasmids and then observed that its overexpression promoted evident changes in the cells morphology (EMT, epithelial to mesenchymal transiton) (Fig. 1B). To confirm that the changes of morphology are induced by EMT, we performed immunoflurescence analysis to detect epithelial and mesenchymal markers of HEC-1A cells transfected with L1CAM expressing plasmids and empty vectors. We found that the E-Cadherin protein (epithelial marker) was inhibited and Vimentin protein (mesenchymal marker) were induced by L1CAM in HEC-1A cells (Fig. 1C). To further analyze whether L1CAM could affect E-Cadherin and Vimentin protein, we used western blotting to detect their expression in the cells transfected with L1CAM expressing plasmids and empty vectors. The results demonstrated that E-Cadherin was downregulated and Vimentin was upregulated by L1CAM (Fig. 1D and E). We also performed real-time PCR to detect epithelial and mesenchymal markers. As anticipated, we found that epithelial marker (E-cadherin) was downregulated and mesenchymal markers (such as N-Cadherin, Vimentin, and ZEB1) was upregulated by L1CAM in HEC-1A cells (Fig. 1F).

LICAM promotes formation of CICs in HEC-1A cells. EMT can contribute to increased formation of CICs in cancer cells (34-37). To determine whether L1CAM could affect characteristics of CICs, we performed sphere forming assay to evaluate the formation of CICs in HEC-1A cells. The results of sphere forming assay showed that formation of spheres were increased by L1CAM in HEC-1A cells (Fig. 2A). Moreover, we performed western blot to detect whether L1CAM could regulate CICs markers-Musashi-1 and CD133 expression in the cells. We found that Musashi-1 and CD133 protein were evidently upregulated by L1CAM in HEC-1A cells (Fig. 2B and C).

*LICAM promotes paclitaxel resistance in human endometrial cancer HEC-1A cells*. To further identify whether L1CAM can affect paclitaxel efficacy in HEC-1A cells, we performed MTT



Figure 1. L1CAM promotes EMT in endometrial cancer HEC-1A cells. (A) Western blot of L1CAM in L1CAM expressing plasmids and empty vectors (mock) transfected HEC-1A cells. n=3. (B) HEC-1A cells transfected as indicated were photographed after 72 h of transfection. n=3. (C) Immunofluorescence assay of E-cadherin and vimentin in L1CAM expressing plasmids and empty vectors (mock) transfected HEC-1A cells. n=3. (D) Western blot of E-cadherin in L1CAM expressing plasmids and empty vectors (mock) transfected HEC-1A cells. n=3. (E) Western blot of vimentin in L1CAM expressing plasmids and empty vectors (mock) transfected HEC-1A cells. n=3. (E) Western blot of vimentin in L1CAM expressing plasmids and empty vectors (mock) transfected HEC-1A cells. n=3.



Figure 2. L1CAM promotes formation of CICs in endometrial cancer HEC-1A cells. (A) Sphere growth of HEC-1A cells transfected with L1CAM expressing plasmids and empty vectors (mock). (B) Western blot of Musashi-1 in L1CAM expressing plasmids and empty vectors (mock) transfected HEC-1A cells. n=3. (C) Western blot of CD133 in L1CAM expressing plasmids and empty vectors (mock) transfected HEC-1A cells. n=3.

assay in HEC-1A cells treated as indicated (Fig. 3A). The results showed that overexpressing L1CAM could promote paclitaxel resistance (Fig. 3A). In addition, we performed western blot to analyze cyclophilin A protein expression in L1CAM expressing plasmids and empty vectors transfected HEC-1A cells. We found that cyclophilin A protein can be increased by L1CAM (Fig. 3B).



Figure 3. L1CAM promotes paclitaxel resistance in endometrial cancer HEC-1A cells. (A) MTT of HEC-1A cells transfected with L1CAM expressing plasmids and empty vectors (mock) were treated with different concentration of paclitaxel. n=3. (B) Western blot of cyclophilin A in L1CAM and empty vectors (mock) transfected HEC-1A cells. n=3.

LICAM regulates paclitaxel resistance-associated microRNAs expression in HEC-1A cells. Oncogenes can promote endometrial cancer progression by regulating microRNA expression and microRNA involved in endometrial cancer pathogenesis can function as oncogene or tumor suppressor gene (38-40). Thus, we reasoned that L1CAM could function as an oncogene by regulating miRNAs expression. We performed microarrays to detect miRNA expression. RNAs isolated from L1CAM expressing plasmids or empty vectors transfected HEC-1A cells were hybridized to a custom miRNA microarray platform. We found that miR-135a, miR-375, miR-200c, miR-182, let-7e, miR-31, miR-21, miR-200b, miR-143 and miR-145 were changed more than 10 folds in the cells (Fig. 4A).

L1CAM promotes anoikis resistance in human endometrial cancer HEC-1A cells. To study the roles of L1CAM on metastasis, we used anoikis assays to detect its role regulating anoikis resistance. Cells transfected with L1CAM expressing plasmids showed about 200% increased resistance to anoikis-mediated cell death (Fig. 4B).

#### Discussion

The expression of L1CAM is a strong predictor of poor outcome in endometrial cancer (41). EMT plays an important role in invasion and metastasis of endometrial cancer and enables cancer cells to obtain malignant characters and traits of CICs (42). Critical molecular features of this process are the deregulation of E-cadherin and vimentin expression (43). Consistent with previous report that L1CAM was inversely associated with E-cadherin expression (44), we found that overexpressing L1CAM induced EMT and inhibited E-cadherin expression in endometrial cancer HEC-1A cells.

CICs have been proposed as the major power of EMT and responsible for poor survival (45). Musashi-1 and CD133 have been proposed as markers of CICs for endometrial cancer (46). In line with previous report that L1CAM is required for



Figure 4. L1CAM regulates paclitaxel resistance-associated microRNAs expression and promotes anoikis resistance in endometrial cancer HEC-1A cells. (A) miRNA microarray analysis for L1CAM expressing plasmids or empty vectors (mock) transfected HEC-1A cells. n=3. (B) Anoikis assays for L1CAM expressing plasmids or empty vectors (mock) transfected HEC-1A cells. n=3.

maintaining CICs and targeting L1CAM may represent a novel therapeutic strategy (47), we showed that its overexpression evidently promoted formation of CICs traits and upregulated Musashi-1 and CD133 protein. All the results indicated that L1CAM might be a therapeutic target for eradicating CICs in endometrial cancer.

Chemotherapy is a common therapeutic strategy for cancer, but it fails to eradicate cancer cells, because of primary resistance or acquired drug resistance. Elucidating the mechanisms of drug resistance for cancer will yield vital information about how to improve cancer chemotherapy and circumvent the resistance. In line with previous report that L1CAM can confer chemoresistance in malignant tumor (48,49), we showed that over-expressing L1CAM could promote paclitaxel resistance in endometrial cancer HEC-1A cells. Cyclophilin A expression was increased in paclitaxel-resistant endometrial cancer cells, as well as silencing Cyclophilin A reversed paclitaxel resistance (50). We showed that Cyclophilin A was upregulated by L1CAM. microRNAs have recently been identified as key genes implicated in mechanisms of chemoresistance. Upregulation of miR-135a and miR-375 can contribute to paclitaxel resistance. Up-regulating miR-200c in ovarian cancer reduced tumor burden and improved paclitaxel sensitivity. We showed that L1CAM significantly upregulatedmiR-135a and miR-375 expression and downregulated miR-200c expression. The results indicated that L1CAM may induce paclitaxel resistance by regulating microRNAs.

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