Bovine endometrial MSC: mesenchymal to epithelial transition during luteolysis and tropism to implantation niche for immunomodulation

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Abstract

Background: The uterus is a histologically dynamic organ, and the mechanisms coordinating its regeneration during the oestrous cycle and implantation are poorly understood. The aim of this study was to isolate, immortalize and characterize bovine endometrial mesenchymal stem cell (eMSC) lines from different oestrous cycle stages (embryo in the oviduct, embryo in the uterus or absence of embryo) and examine their migratory and immunomodulatory properties in an inflammatory or implantation-like environment, as well as possible changes in cell transdifferentiation.

Methods: eMSCs were isolated and analysed in terms of morphological features, expression of cell surface and intracellular markers of pluripotency, immunocytochemical analyses, alkaline phosphatase activity, proliferation and osteogenic or chondrogenic differentiation capacities, as well as their ability to migrate in response to inflammatory (TNF-α or IL-1β) or implantation (IFN-τ) cytokines and their immunomodulatory effect in the proliferation of T cells.

Results: All eMSCs showed MSC properties such as adherence to plastic, high proliferative capacity, expression of CD44 and vimentin, undetectable expression of CD34 or MHCII, positivity for Pou5F1 and alkaline phosphatase activity. In the absence of an embryo, eMSC showed an apparent mesenchymal to epithelial transition state. eMSC during the entire oestrous cycle differentiated to osteogenic or chondrogenic lineages, showed the ability to migrate in response to inflammatory (TNF-α or IL-1β) or implantation (IFN-τ) cytokines and their immunomodulatory effect in the proliferation of T cells.

Conclusion: This study describes for the first time the isolation, immortalization and characterization of bovine mesenchymal stem cell lines from different oestrous cycle stages, with a clear mesenchymal pattern and immunomodulatory properties. Our study also reports that the migratory capacity of the eMSC was increased towards an inflammatory niche but was reduced in response to the expression of implantation cytokine by the embryo. The combination of both signals (pro-inflammatory and implantation) would ensure the retention of eMSC in case of pregnancy, to ensure the immunomodulation necessary in the mother for embryo survival. In addition, in the absence of an embryo, eMSC showed an apparent mesenchymal to epithelial transition state.

Keywords: Endometrial mesenchymal stem cells, Embryo implantation, Inflammation, Cell migration

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Background
In ruminants, embryo mortality is a key factor affecting fertility [1]. Approximately 50% of pregnancy loss occurs between blastocyst hatching and embryo attachment to endometrium in cattle [2]. Efficient communication between cells and tissues is paramount in many physiological processes, including embryo development. Preimplantational embryos of mammals develop within the female genital tract (oviduct and uterus) and their efficient communication conditions their development and survival. In ruminants, the bicornate uterus is coated with the endometrium, the ultimate and unique biological layer facing the implanting embryo in normal early pregnancy, which drives the development of the embryonic disc and extra-embryonic tissues [3]. The endometrium displays two specific areas, namely the caruncles and the intercaruncular areas. The caruncles represent aglandular structures of limited size that are distributed over the endometrial surface. The intercaruncular areas are large and contain the endometrial glands, which produce histotroph, a collection of numerous and diverse factors including cytokines and growth factors [4].

During the window of implantation, after blastocyst attach onto the endometrium, both in human and mice, characterized by haemochorial implantation, the stromal cells at the implantation site begin to undergo decidualization, a process by which the fibroblast-like endometrial stromal cells differentiate into polygonal epithelial-like cells [5]. The differentiation of decidual cells is one of the earliest uterine adaptations to pregnancy in these species [6] and involves cell growth, change of shape, multinucleation and establishment of inter-cellular junctions [7]. Many genes and signal pathways are involved in the process of decidualization that is crucial for blastocyst implantation and maintenance of pregnancy. Ruminants and pigs have noninvasive implantation; however, they differ in the type of placentaion. Pig conceptuses undergo epitheliochorial placentaion in which luminal epithelium (LE) remains morphologically intact throughout pregnancy and the conceptus trophodeerm simply attaches to the apical LE surface without contacting uterine stromal cells [8]. In contrast, synepitheliochorial placentaion in ruminants involves extensive erosion of the LE due to syncytium formation with trophodeerm binucleate cells. After day 19 of pregnancy, the conceptus tissue is apposed to, but does not penetrate, uterine stroma. Johnson el al. reported that the uterine stroma of sheep undergoes a programme of differentiation similar to decidualization in invasive implanting species, whereas porcine stroma exhibits differentiation that is more limited than that in sheep, rodents or primates [9].

The uterus is a histologically dynamic organ, and the mechanisms coordinating its regeneration during the oestrous cycle and implantation are poorly understood. It has been proposed that bone marrow-derived cells contribute to uterine regeneration [10]. An alternative or additive mechanism of endometrial regeneration could be in situ cellular transdifferentiation, an example of which is mesenchymal to epithelial transition (MET). During MET, mesenchymal cells are re-programmed, thereby gradually losing mesenchymal cell characteristics while gaining epithelial cell traits [11]. MET and epithelial to mesenchymal transition (EMT) are fundamental processes that occur during the bovine embryo development and are also implicated in tumour metastasis [12]. Uchida et al. reported that the EMT might play an important role in human embryo implantation helping the embryo to penetrate through the endometrial epithelial cells into the endometrial stromal cell layer [13]. In addition, Zhang et al. suggested that MET might exist during decidualization, providing a stable developmental environment and an anchor point for embryos to invade into the uterus [14]. Recently, Patterson et al. reported that MET contributes to endometrial regeneration following natural and artificial decidualization by sesame oil injection into the uterine lumen in mouse [15], but the possibility that this cellular transdifferentiation also serves as a mechanism to regenerate adult endometrium in bovine has not been evaluated.

Although the presence of stem cells in the endometrium was first speculated in 1978 by Priianishnikov [16], it was not until 2004 that stem cells were identified and characterized for the first time in the endometrial tissue [17, 18]. The adult stem cells, mesenchymal stem cells (MSCs), unlike pluripotent embryonic stem cells, are multipotent and can produce a limited range of cell lineages depending on their location. MSCs are isolated from virtually every tissue, so that their stem cell properties are difficult to predict in an in vivo setting. The presence in the bovine endometrium of mesenchymal stem cells (eMSCs) has been already described [19–21]. Bovine eMSCs possess several MSC properties such as clonogenicity, ability to adhere to plastic, fibroblast morphology and in vitro differentiation capacity to adipocyte, chondrocyte and osteocyte [22]. Moreover, bovine eMSCs present high levels of CD29, CD44 and vimentin markers, low expression levels for CD34 and no MHC-II [21]. It is important to highlight that in some cases MSCs are able to divide but to a limited extent in vitro before entering replicative senescence. Between passages 7 and 12, MSCs increase their cell size and reduce the expression of certain pluripotency markers, leading to proliferative arrest [23].

MSCs have the ability to migrate to damaged and inflamed tissues where they proliferate and utilize their immunomodulatory properties as part of the regeneration process of the tissue itself [24–26]. The immune
reaction has classically been functionally divided into Th-1 and Th-2 responses. Although, in human, numerous studies attempted to support pregnancy as a Th-2 or anti-inflammatory state [27], a similar number of studies supported its condition as a Th-1 or pro-inflammatory state [28]. A two-phase model was thereafter proposed by Mor et al., being the first and third trimesters characterized by a pro-inflammatory ambient, while the second trimester would represent an anti-inflammatory phase [29]. MSC can suppress or reduce the proliferation of T cells and also have a role in the regulation of Th-1/Th-2 balance shifting a Th-1 phenotype to a Th-2 [30]. MSC interactions with immune cells have been studied widely in both humans and rodents even in pigs [24], although this research has not yet been extended to MSC from the main veterinary species.

In ruminants, the major embryo-derived signal of pregnancy recognition has been suggested to be type I interferon [31]. The production of this interferon (interferon-tau; IFN-τ) is unique as it is secreted by trophoderm cells during the elongation phase and secretion stops when trophoderm cells appose to the endometrial luminal epithelium. IFN-τ is crucial for maternal recognition through the inhibition of endometrial prostaglandin-F2 alpha secretion, thus preventing luteolysis, an indispensable step for maintaining P4 luteal secretion [32].

Taking all this information into account, we have addressed the isolation, immortalization and characterization of eMSC from different phases of the bovine oestrous cycle in terms of their ability to migrate in vitro by agarose spot assay mediated by inflammatory cytokines (TNF-α or IL-1β) or the implantation cytokine (IFN-τ). We have also evaluated their possible immunomodulating function in a xenogeneic condition.

**Methods**

**Isolation and culture of bovine endometrial mesenchymal stem cells (eMSCs)**

Heifer uterus and ovaries were collected from routine slaughtered at the local abattoir. Oestrous cycle stages were determined by a macroscopic examination of the luteum corpus and ovarian morphology of the cycling heifers according to literature criteria [33] (see Table 1). The uterine horns ipsilateral to the luteum corpus were removed and washed with fresh tap water, and the uterine lumen was washed three times using a 20-ml syringe with Hank’s balanced salt solution (HBSS) supplemented with 500 U/ml penicillin, 500 mg/ml streptomycin and 0.1% bovine serum albumin (BSA) (Merck KGaA, Darmstadt, Germany) by flushing from the utero-tubal junction. To obtain bovine endometrial cells, a longitudinally cut of the uterine horn was performed with sterile scissors. The intercaruncular endometrium was carefully dissected from underlying tissues, cut in strips, minced using sterile scissors and incubated in 25-ml type II collagenase solution: HBSS supplemented 0.05% type II collagenase (Gibco by Life Technologies, Grand Island, NY, USA), 0.1% BSA (Merck KGaA, Darmstadt, Germany) and 30 mM CaCl2. The solution was homogenized and incubated during 1 h at 37 °C shaking gently every 10 min. The solution resulting from the digestion was filtered through a 70-μm cell strainer (SPL life science, Korea), and a volume of culture medium RPMI (Hyclone Laboratories, Utah, USA), supplemented with 10% foetal calf serum (PAA laboratories, Austria), 2% non-essential amino acids and antibiotics (100 U/ml penicillin, 100 mg/ml streptomycin), was added to block the action of collagenase and the obtained suspension centrifuged at 300 × g for 5 min. The resulting pellets were resuspended in culture medium and plated in 100-mm² tissue culture dish (JetBiofil, Guangzhou, China) and incubated in an atmosphere of humidified air and 5% CO2 at 37 °C. Culture medium was changed every 48–72 h.

**eMSC immortalization**

PA-317 LXSN-16E6E7 cells were acquired from the American Type Culture Collection (Manassas, VA). Virus was harvested from 16-h conditioned media of PA-317 cells that and filtered through a 0.45-mm filter (Millipore, Bedford, MA) to remove cells and debris. Ten microlitres of viral supernatant was diluted 1:3 with serum-free DMEM/RPMI medium containing polybrene (4 mg/ml; Sigma). Primary cultures of bovine endometrial cells obtained from different cyclic stages which have been grown to 60% confluence in P100 culture dishes were overlaid

| Oestrous cycle stage   | Embryo location                                      | Primary cultures | Immortalization efficiency | Immortalized eMSC | Oestrous stage |
|------------------------|-----------------------------------------------------|------------------|-----------------------------|-------------------|---------------|
| D2-D4 early luteal     | Embryo in the oviduct                               | 4                | 1 (25%)                     | 1A                | 1             |
| D4-D8 medium luteal    | Embryo in the uterus with zona pellucid (ZP)        | 4                | 0 (0%)                      | –                 | 2             |
| D9-D16 late luteal     | Embryo in the uterus w/o ZP                         | 8                | 3 (37.5%)                   | 3A, 3D, 3E        | 3             |
| D17-D21 follicular phase | Regression of corpus luteum if not embryo         | 6                | 4 (66.7%)                   | 4B, 4C, 4D, 4H   | 4             |
with 3 ml of this mixture and incubated in an atmosphere of humidified air and 5% CO2 at 37 °C for 3 h. After 3–4 h, viral supernatant was removed; cells were washed twice with HBSS, and complete culture medium was added to dishes. After 48 h, cells having integrated the vector were selected by resistance to the neomycin analogue G418 300 μg/ml (Gibco of Life Technologies, Grand Island, NY, USA) in complete culture medium. Selection continued for 7 days, and surviving cells were subsequently cultured with complete culture medium containing 100 μg/ml G418 for general cell maintenance. The G418-resistant cell populations were subcultured for more than 30 passages, and immortalized cells were cryopreserved each passage.

**eMSC characterization by flow cytometry**

Surface, cytoplasmic and nuclear cell antigens were analysed by flow cytometry using a Cell Lab Quanta SC system of Beckman Coulter. Cell cultures at 80–90% confluence were detached using 0.05% trypsin–EDTA (T/E) solution, collected and fixed with 4% paraformaldehyde for 10 min and subsequently washed twice with PBS. For the analysis of the expression of vimentin (clone LN-6, Sigma-Aldrich) and cytokeratin (Clone C-11, Sigma-Aldrich) (cytoplasmic proteins) and POU5F1 (rabbit polyclonal, Biorbit) (a nuclear protein), cell permeabilization was performed by incubation with 0.3–0.5% Triton X-100 for 10 min and washing with PBS. Non-specific binding of antibodies was blocked with TNB-blocking solution during 30 min at 37 °C. Appropriate dilutions, provided by manufacturers, of primary antibodies against the markers commonly used to define mesenchymal stromal cells (MSCs): vimentin (clone LN-6, Sigma-Aldrich) and CD44 (clone IM7, Bio-rad), as positive markers; cytokeratin, CD34 (Rabbit polyclonal, Biorbit), and MHCII (clone CVS20, Bio-Rad) as negative markers; and POU5F1 as pluripotency marker, were added to the cells and incubated overnight at 4 °C. Cells were then stained with the appropriate Alexa Fluor 488-conjugated secondary antibodies (Jackson ImmunoResearch Laboratories, Baltimore Pike, West Grove, PA). Negative control samples were produced by the omission of the primary antibody. The analysis of the samples was performed with Cell Lab Quanta SC system of Beckman Coulter using Flow-Jo X SOFTWARE version 10.0.7r2.

**Alkaline phosphatase (AP) activity**
eMSC lines at passages 10–15 were grown on 35-mm dishes (JetBiofil, Guangzhou, China) for 2 weeks. Cells were washed twice with PBS and fixed with a solution of 4% paraformaldehyde for 10 min at room temperature. Paraformaldehyde was aspirated, and the plates were washed twice with distilled water and covered with B solution (1 ml of Solution A (Fast Red 1 mg/ml), 1.6 μl of Naphtol AS-mx phosphate and 40 μl TRIS HCl 1 M, pH 8.6) during 10–15 min at room temperature in the dark. Solution B was finally removed, and the cells were washed twice with PBS and covered with PBS to prevent drying. The colonies were studied for appearance of pink/red coloration indicating alkaline phosphatase activity. The stained colonies were imaged using a Motic SMZ-171 stereomicroscope coupled to a Moticam BTU8 digital camera using Motic Image Plus software version 2.0 (Motic China Group Co., LTD).

**Cell proliferation measurement**

Different mesenchymal cell lines of passages 10–15 were seeded at 2 × 105 cells per 60-mm tissue culture plates (JetBiofil, Guangzhou, China). The culture medium was changed every 2 days. At each time point, a duplicate of plates was detached by tripnisization and counted using a Bürker counting chamber (Paul Marienfeld GmbH & Co. Lauda-Königshofen, Germany). Twenty microlitres of cell suspension was placed on both sides of the chamber and viewed using ×100 magnification under an inverted Nikon Diaphot phase contrast microscope coupled to a Jenoptik ProRes CT1 digital camera. Images were taken using ProRes capture pro software version 2.7 (Jenoptik Laser, Optic Systeme GmbH). A total of 5 × 1 mm² squares per sample were analysed, and the number of cells per millilitre was determined according to number of cells in 1 ml = N/Z × dilution × 10⁴, where N = the whole number of cells counted and Z = the number of counted squares. Cell population doubling time (PDT) was calculated using Roth V. 2006 Doubling Time Computing, available from http://www.doubling-time.com/locate.php.

**In vitro differentiation potential assay**
eMSC lines at passages 10–15 were grown until 90% confluence on 24-well multidish (JetBiofil, Guangzhou, China). For osteogenic differentiation, The StemPro® Osteogenesis Differentiation Kit (Thermo Fisher Scientific, Meridian Road Rockford, IL, USA) was used according to the manufacturer’s instructions. Differentiating media was changed every 3–4 days for 21 days. Simultaneously, control cells were cultured in standard conditions. Cells were then fixed in 4% paraformaldehyde solution for 30 min. After fixation, cells were incubated for 2–3 min in 2% Alizarin Red S solution (pH 4.2) to visualize the calcium deposits.

For chondrogenic differentiation, The StemPro® Chondrogenesis Differentiation Kit (Thermo Fisher Scientific, Meridian Road Rockford, IL, USA) was used according to the manufacturer’s instructions. Differentiating media was changed every 2–3 days for 14 days. Simultaneously, control cells were cultured in standard conditions. Cells
were then fixed in 4% paraformaldehyde solution for 30 min. After fixation, cells were incubated for 30 min with 1% Alcian Blue solution prepared in 0.1 N HCl. Blue staining was corresponding with proteoglycans synthetized by chondrocytes. Cells under differentiation conditions were photographed, using a Motic SMZ-171 stereomicroscope coupled to a Moticam BTU8 digital camera.

**Cell migration measurement: agarose spot assay**

The cell migration measurement by agarose spot assay was carried out following the procedures of Calle et al. [34]. Briefly, PBS-0.5% agarose solution was heated on a water bath until boiling to facilitate complete dissolution. When the temperature cooled down to 40 °C, 90 µL of agarose solution was pipetted into a 1.5-ml Eppendorf tube containing 10 µL of PBS or PBS supplemented with TNF-α, IL-1β or IFN-γ for a final concentration of 6 nM [33]. Five-microlitre spots of agarose-containing PBS, TNF-α or IL-1β were pipetted onto six-well plates (JetBiofil, Guangzhou, China), 16 drops per well, 12 drops per MSC line, and allowed to cool for 15 min at 4 °C. At this point, cells that had been treated with 1 µg/ml C-Mitomycin overnight (Merck KGaA, Darmstadt, Germany) to avoid cellular duplication were separated using the EasySep magnet (StemCell Technologies). Enriched T cells were then washed and resuspended using Bürker counting chamber. Imaging was performed at 24 and 48 h using a Motic SMZ-171 stereomicroscope coupled to a Moticam BTU8 digital camera using Motic Image Plus software version 2.0 (Motic China Group Co., LTD). Motile cells penetrated the agarose spot. The longest straight distance from the border of the spot was analysed for each cell using ImageJ.

**Statistical analysis**

Statistical analysis was performed using GraphPad Prism 6 (GraphPad Software, La Jolla, CA, USA). One-way ANOVA for multiple comparisons by Fisher’s LSD tests were used for cell proliferation and doubling time. Two-way ANOVA for multiple comparisons by Fisher’s LSD tests were used for cell migration. Values are presented in mean ± standard error of the mean (SEM). Differences were considered to be significant when \( p < 0.05 \). Two-tailed Student’s \( t \) tests were used when two groups were compared for immunomodulatory assays. Values are expressed as mean ± standard error of the mean (SEM). Differences were considered to be significant when \( p < 0.05 \).

**Results**

**MSC isolation and immortalization efficiency**

eMSCs were isolated from heifer uterus that were ascribed to one of the four bovine oestrus stage categories (1–4) (Table 1) proposed by [33] based on the morphology of the active ipsilateral ovary to the uterine horn from which the cells were isolated. To avoid the risk of senescence by the maintenance of eMSC in vitro, isolated cells were immortalized using the retroviral vector LXSN-16E6E7. From a total of 22 primary cultures of endometrial stromal cells, eight cell lines were successfully immortalized (eMSC-1A, eMSC-3A, eMSC-3D, eMSC-3E, eMSC-4B, eMSC-4C, eMSC-4D, eMSC-4H) (Table 1).

**Morphological features**

Pre-immortalized mesenchymal stem cell cultures at passage 0 adhered to the plastic surface of culture dishes exhibiting a mixture of round, spindle or elongated shape morphology (Fig. 1—upper panels). However, after the first cell passage, the cells formed a more homogeneous population of fibroblast-like adherent cells, with the exception of eMSC-4D and eMSC-4H that showed an epithelial-like morphology remained constant before and after the immortalization even after more than 20 passages (Fig. 1—lower panels).
Expression of cell surface, intracellular and pluripotent-specific markers

Some characteristic MSC surface and intracellular markers were assessed by flow cytometry (Fig. 2a–c). All cell lines were positive for cell surface CD44 and cytoplasmic vimentin, both of them are characteristic markers of MSCs. Interestingly, cytokeratin, a typical cytoplasmic marker expressed by epithelium of ectoderm and endoderm, and commonly used as a negative marker of mesenchymal stem cells, was present in all stage 4 eMSC lines: strongly detected in eMSC-4H, clearly positive in eMSC-4C and slightly positive in eMSC-4B and eMSC-4D (Fig. 2b), correlating with the epithelial morphology of two of these cell lines. No expression of haematopoietic markers, such as MHCII or CD34, was found in any of the eMSC lines. Regarding pluripotency features, all eMSC lines, including those cell lines immortalized from the follicular phase and with epithelial morphology, were positive for the nuclear marker POU5F1 (Fig. 2c) and showed alkaline phosphatase activity (Fig. 2d).

Proliferation capacity

To analyse the cell proliferation capacity of eMSCs, the number of cells/dish was counted for each cell line at days 3, 5, 7 and 10, starting in all cases from an initial seeding of $2 \times 10^5$ cells. As showed in Fig. 3a, the number of cells increased for all cell lines along the entire assay, demonstrating high proliferative capacity of these cells even at passages 10–15. eMSC-4H was the most proliferative line and at day 10 reached $19.74 \times 10^5 \pm 2.13 \times 10^5$ cells. Figure 3b shows the proliferation rate of MSC lines in terms of the population doubling time (PDT). Among eMSC lines, eMSC-4H and eMSC-4D
showed on day 10 a significantly shorter PDT of 5.77 ± 0.47 and 3.50 ± 0.23 days, respectively ($p < 0.05$).

**In vitro differentiation of MSCs**

As showed in Fig. 4, all eMSC lines cultured under osteogenic conditions presented a comparable amount of calcium deposits, indicating a high osteogenic differentiation potential of these lines. Cells cultured under chondrogenic conditions showed the presence of acidic proteoglycan that was demonstrated at monolayer cells by Alcian blue staining, presenting in most cases stained nodules typical from cartilaginous tissue.

All this characterization demonstrates that all immortalized lines complied with MSC features in terms of marker expression and differentiation capacities, even those obtained from the follicular phase that showed cytokeratin expression and epithelial morphology.

**Migratory response of MSC to inflammatory and implantational stimuli**

Assessment of the invasive capacity of all eMSC lines was performed using the agarose spot assay [34], which allows the quantification of cell invasion by measuring their crawling underneath an agarose gel on a planar surface. All eMSC lines showed migration capacity into agarose spots without cytokine stimuli from day 2 (Fig. 5). In terms of migration without cytokine stimuli, eMSC-4C showed a significantly higher migration rate than the rest of eMSC lines with a net migrated distance of 1008.0 ± 38.26 μm on day 7 (Fig. 5B).

The inflammatory cytokine TNF-α did not cause an increase in the migration capacity of any of the eMSC lines studied compared to the control of PBS, but completely abrogated that of eMSC-3D and eMSC-4B. In contrast, the inflammatory cytokine IL-1β stimulated the

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**Fig. 3** In vitro proliferation of mesenchymal stem cells. **a** Absolute number of cells/dish (mean ± SD) at indicated points. **b** Doubling time in days of each MSC line (mean ± SD). Different letters indicate significant differences ($p < 0.05$).

**Fig. 4** In vitro differentiation of eMSC to different lineages. Images show Alizarin Red S staining of calcium deposits in cells cultured in basal medium (control) or in osteogenic differentiation medium (top panel); Alcian blue staining of acidic proteoglycan in cells cultured in basal medium (control) or in chondrogenic differentiation medium (bottom panel). Bright field images were acquired with a ×3 magnification (bars = 150 μm).
migration capacity of cell lines eMSC-1A, eMSC-3A, eMSC-3E, eMSC-4C and eMSC-4H while caused a reduction in the cell migration rate of only the eMSC-4B line. Regarding the preimplantation protein IFN-\(\tau\), it reduced cell motility of most cell lines (Fig. 5C).

**Immunomodulatory properties**

Although all eMSC immortalized lines showed MSC differentiation capacities, those obtained from the follicular phase had unexpected features such as cytokeratin expression and epithelial morphology. Thus, we analysed the immunomodulatory potential of two representative lines: eMSC-1A with a pure mesenchymal pattern and eMSC-4H with a phenotype consistent with MET. Co-culture experiments with human T cells activated with anti-CD2/3/28 beads were established to analyse eMSC's influence on polyclonal xenogeneic proliferation. The percentage of proliferating cells was determined by flow cytometry, quantifying FSC\(^{\text{high/Violet}^{\text{low}}}\) T cell blasts (Fig. 6a, b). An almost complete abrogation of activated human T cell proliferation was observed when these cells were co-cultured with either eMSC-1A (\(p < 0.0001\)) or eMSC-4H (\(p < 0.0001\)) (Fig. 6c).

**Discussion**

We have successfully isolated, immortalized and characterized eMSC from different phases of the bovine oestrous cycle. The results of the present study clearly demonstrated that immortalized eMSC showed MSC properties such as adherence to plastic, high proliferative capacity until passages 10–15 and surface expression of CD44 and intracellular expression of vimentin, both MSC markers. Moreover, all eMSC lines were negative...
for haematopoietic specific markers CD34 and MHCII and positive for pluripotency-related marker Pou5F1, a nuclear protein that is well known for pluripotency maintenance and is expressed in bovine MSCs [20]. Numerous studies demonstrate that the ability to express alkaline phosphatase activity is also a pluripotency marker of stem cells [35]. In this report, immortalized eMSC lines from different phases of the bovine oestrous cycle showed positivity in AP activity. All eMSC lines from early and late luteal phases showed a fibroblast-like morphology characteristic of MSC and expressed vimentin. Interestingly, eMSC lines obtained from the follicular phase (absence of an embryo) expressed cytokeratin and two of these lines showed a clear epithelial-like morphology. This observation could reflect a mesenchymal to epithelial transition (MET) of eMSC, a process in which mesenchymal cells are re-programmed, thereby gradually losing mesenchymal cell characteristics while gaining epithelial cell traits [11]. In both mouse and human, uterine regeneration was shown to occur both by bone marrow-derived cells [10] and also by MET of resident cells [14, 15]. Cousins et al. demonstrated that MET contributes to endometrial epithelium after progesterone withdrawal-induced tissue breakdown in a mouse model of menstruation [36]. This finding is consistent with Yu et al. observations showing that luteal hormone concentration and/or duration thresholds confer plasticity and reversibility of endometrial stromal phenotype via MET [37]. In bovine, progesterone produced by the corpus luteum is essential for the establishment and maintenance of pregnancy. Rapid regression of the cow’s corpus luteum is a key event in the bovine oestrous cycle as it is responsible for the precipitous decrease in the blood concentration of progesterone, the end of the cycle (follicular phase) [38]. Our eMSC lines established from the different phases of the oestrous cycle show a clear mesenchymal pattern in the early luteal phase (eMSC-1A, corresponding to the
presence of the embryo in the oviduct) and late luteal phase (eMSC-3A, eMSC-3D and eMSC-3E, corresponding to the presence of the embryo in the uterus), while an apparent mesenchymal to epithelial transition state is observed in eMSC from the follicular phase (eMSC-4B, eMSC-4C and eMSC-4D, and eMSC-4H corresponding to the regression of the corpus luteum if there was no embryo) since they express cytokeratin and in some cases show an epithelial morphology. Considering that both the epithelium and the bovine stroma present oestrogen and progesterone receptors during the oestrous cycle [39], it would be interesting to analyse in future experiments the paracrine function performed by the eMSCs, thanks to the possible expression of hormone receptors during the oestrous cycle.

The ability to differentiate in vitro to mesodermal lineages has been demonstrated for eMSC isolated from the endometrium at the medium luteal phase [20] or medium and late luteal phase [21], but it was not known if the MSCs present in the uterus conserved such multipotency capacity during the entire oestrous cycle. Our data on osteogenic and chondrogenic induction with eMSC lines confirmed their multipotency, both in the early and late luteal phases as well as in the follicular phase.

An important characteristic of MSCs is their ability to migrate to damaged tissues or to inflammatory processes. All eMSC lines showed migration capacity without cytokine stimulation. There was not a clear difference in terms of non-stimulated migration between eMSC lines from different oestrus phases or with different morphologies (mesenchymal or epithelial). Although the eMSC lines obtained from the follicular phase (absence of an embryo) expressed cytokeratin and two of these lines showed clear epithelial type morphology, they also showed a high migration capacity and showed intracellular expression of vimentin. Vimentin is an important marker of mesenchymal to epithelial transition and a requisite regulator of mesenchymal cell migration. During MET, vimentin protein and mRNA expression are downregulated as cell motility decreases and cells adopt epithelial characteristics [40, 41]. Taking into account the above, MSCs show a cellular plasticity characteristic of MET that allows them to show characteristics of mesenchymal and epithelial cells. In all scenarios, either when the embryo is still in the oviduct (phase 1), or when the embryo without zona pellucida is in the uterus (phase 3), or in case of regression of the corpus luteum as a consequence of the absence of an embryo in the uterus (phase 4), most of the eMSCs present an active migration towards a pro-inflammatory niche (Th-1) characterized by the presence of IL-1β and described in the initial and final stages of human gestation [29]. In contrast, most of the eMSCs from the three scenarios described respond with a block in their migration to IFN-τ, the major embryo-derived pregnancy signal in bovine [31]. This is in clear contrast with the effect of this cytokine on bovine MSC derived from bone marrow, which are attracted by IFN-τ (Calle et al., under revision). The combination of both IL-1 and IFN-τ signals would ensure the retention of eMSC in case of pregnancy, as well as further recruitment of MSC from the bloodstream, to ensure the
immunomodulation necessary in the mother for embryo survival.

eMSCs could act there as primarily suppressive effectors on Th1 cells by suppressing their proliferation, differentiation and activation [42, 43]. In accordance, both analysed lines; eMSC-1A, with a pure mesenchymal pattern and eMSC-4H1 with a phenotype consistent with MET, showed the ability to suppress T cell proliferation in vitro.

Conclusion
In sum, our results show that eMSC isolated and immortalized from the uterus shared common MSC features, although eMSCs isolated from the endometrium of the final phase of the oestrous cycle show a phenotype compatible with the mesenchymal-epithelium transition. Finally, the migratory capacity of eMSCs towards an inflammatory niche and their retention by interferon-γ secreted by the embryo would ensure the immunomodulatory ambient necessary for productive pregnancy (Fig. 7).

Abbreviations
BSA: Bovine serum albumin; DMEM-LG: Dulbecco’s modified Eagle’s medium low glucose; eMSCs: Endometrial mesenchymal stem cells; EMT: Epithelial to mesenchymal transition; HBSS: Hank’s balanced salt solution; IFN-γ: Interferon-γ; IL-1β: Interleukin-1β; LE: Luminal epithelium; MET: Mesenchymal to epithelial transition; MHCII: Major histocompatibility complex II; PDT: Cell population doubling time; POU5F1: POU class 5 homeobox 1; T/E: Trypsin−EDTA; TAC: Tetrameric antibody complexes; TNF-α: Tumour necrosis factor-α.

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Availability of data and materials
The materials used and/or analysed during the current study are available from the corresponding author on reasonable request.

Authors’ contributions
MAR contributed to the conception and design and financial support. AC and MAR contributed to the provision of study material, collection and assembly of data, data analysis and interpretation, manuscript writing and final approval of manuscript; SLM contributed to the provision of study material and critical review of manuscript; BFE contributed to the data interpretation and critical review of manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate
Whole blood, used as source of T lymphocytes for in vitro allogeneic proliferation experiments, was obtained from healthy donors after informed consent approved by the Germans Trias i Pujol University Hospital Ethics Committee. This article does not include any individual patient’s data.

Heifer uterus and ovaries were collected from routine slaughtered at the local abattoir.

Consent for publication
Not applicable.

Competing interests
The authors declare that they have no competing interests.

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References
1. Wiltbank MC, Baez GM, Garcia-Guerra A, Toledo MZ, Monterio PL, Melo LF, et al. Pivotal periods for pregnancy loss during the first trimester of gestation in lactating dairy cows. Theriogenology. 2016;86:239–53.
2. Wolf E, Arnold GJ, Bauersachs S, Beier HAA, Blum H, Einspanier R, et al. Embryo-maternal communication in bovine - strategies for deciphering a complex cross-talk. Reprod Domest Anim. 2003;38:726–89.
3. Hue I, Degrelle SA, Campion E, Renard JP. Gene expression in elongating and gastrulating embryos from ruminants. Soc Reprod Fertil Suppl. 2007;64:365–77.
4. Spencer TE, Bazer FW. Uterine and placental factors regulating conceptus growth in domestic animals. J Anim Sci. 2004;82(E-Suppl):E4–13.
5. Dunn CL, Kelly RW, Crichtley HO. Decidualization of the human endometrial stromal cell: an enigmatic transformation. Reprod BioMed Online. 2003;7:151–61.
6. Alam SM, Konno T, Dai G, Lu L, Wang D, Dunmore JH, et al. A uterine decidual cell cytokine ensures pregnancy-dependent adaptations to a physiological stressor. Development. 2007;134:607–15.
7. Li Q, Kanain A, Wang W, Demayo FJ, Taylor RN, Bagchi MK, et al. Bone morphogenetic protein 2 functions via a conserved signaling pathway involving Wnt4 to regulate uterine decidualization in the mouse and the human. J Biol Chem. 2007;282:31725–32.
8. Björkman N. Fine structure of the fetal-maternal area of exchange in the epithelial-mesenchymal and endothelial-chorial types of placentation. Acta Anat Suppl (Basel). 1973;61:1–22.
9. Johnson GA, Burghardt RC, Joyce MM, Spencer TE, Bazer FW, Pfarrer C, et al. Osteopontin expression in uterine stroma indicates a decidualization-like differentiation during ovine pregnancy. Biol Reprod. 2003;68:1951–8.
10. Du H, Taylor HS. Contribution of bone marrow-derived stem cells to endometrium and endometriosis. Stem Cells. 2007;25:2082–6.
11. Hugo H, Ackland ML, Bick T, Lawrance MG, Clements JA, Williams ED, et al. Epithelial–mesenchymal and mesenchymal–epithelial transitions in carcinoma progression. J Cell Physiol. 2007;213:374–83.
12. Imakawa K, Bai R, Kusama K. Integration of molecules to construct the processes of conceptus implantation to the maternal endometrium. J Anim Sci. 2018;96:3009–21.
13. Uchida H, Maruyama T, Nishikawa-Uchida S, Oda H, Miyazaki K, Yamasaki A, et al. Studies using an in vitro model show evidence of involvement of epithelial-mesenchymal transition of human endometrial epithelial cells in human embryo implantation. J Biol Chem. 2012;287:4441–50.
14. Zhang XH, Liang X, Liang XH, Wang TS, Qi QR, Deng WB, et al. The mesenchymal-epithelial transition contributes to endometrial regeneration following natural and artificial decidualization. Stem Cells Dev. 2013;22:2964–74.
15. Pianishnikov VA. On the concept of stem cell and a model of functional-morphological structure of the endometrium. Contraception. 1978;18:213–23.
