




Oviductal glycoprotein 1 (OVGP1) is expressed by endometrial epithelium that regulates receptivity and trophoblast adhesion

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Abstract

Purpose To study the regulation and functions of oviductal glycoprotein 1 (*OVGP1*) in endometrial epithelial cells.

Methods Expression of *OVGP1* in mouse endometrium during pregnancy and in the endometrial epithelial cell line (Ishikawa) was studied by immunofluorescence, Western blotting, and RT-PCR. Regulation of *OVGP1* in response to ovarian steroids and human chorionic gonadotropin (hCG) was studied by real-time RT-PCR. *OVGP1* expression was knockdown in Ishikawa cells by shRNA, and expression of receptivity associated genes was studied by real-time RT-PCR. Adhesion of trophoblast cell line (JAR) was studied by *in vitro* adhesion assays.

Results *OVGP1* was localized exclusively in the luminal epithelial cells of mouse endometrium at the time of embryo implantation. Along with estrogen and progesterone, hCG induced the expression of *OVGP1* in Ishikawa cells. Knockdown of *OVGP1* in Ishikawa cells reduced mRNA expression of *ITGAV*, *ITGB3*, *ITGA5*, *HOXA10*, *LIF*, and *IL15*; it increased the expression of *HOXA11*, *MMP9*, *TIMP1*, and *TIMP3*. Supernatants derived from *OVGP1* knockdown Ishikawa cells reduced the adhesiveness of JAR cells *in vitro*. Expression of *OVGP1* mRNA was found to be significantly lowered in the endometrium of women with recurrent implantation failure.

Conclusion *OVGP1* is specifically induced in the luminal epithelium at the time of embryo implantation where it regulates receptivity-related genes and aids in trophoblast adhesion.

Keywords Endometrium · Implantation · Receptivity · Oviductal glycoprotein 1 · Glycoprotein · Trophoblast

Introduction

Since its first clinical report, assisted reproductive technologies have undergone several advances; however, the success rates remain low. The failure of implantation of morphologically, developmentally, and genetically competent embryos yet remains a major challenge in the success of assisted reproduction

[1, 2]. A significant proportion of women also serially experience implantation failure, and the cause in most of them is unknown [1]. This is because there are gaps in our understanding of fundamental processes of embryo implantation.

Successful implantation involves a receptive endometrium, a developmentally competent embryo, and a synchronized dialogue between maternal and embryonic tissues [3–5]. Even though the blastocyst can implant in different tissues, in the endometrium, it can do so only during a narrow time frame termed as window of implantation or receptive phase [6, 7]. Achieved under the influence of ovarian steroids, the molecular signature of the receptive-stage endometrium is well defined [8, 9]. It is believed that failure to attain receptivity or inability to synchronize embryo transfer with receptive endometrium is a major cause of recurrent implantation failure in women undergoing assisted reproduction [10, 11].

According to the current notion, a receptive endometrium is a passive tissue which readily promotes embryo implantation. However, notwithstanding this notion, we and others have demonstrated an extensive remodeling of the endometrium

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by the embryo at the time of implantation [3, 12–15]. It has been shown [16, 17] that the endometrial remodeling at the time of implantation is mediated by embryo-derived soluble factors like chorionic gonadotropin (CG) and interleukin 1 β (IL1 β). Both in presence of embryo and in response to the soluble factors, the endometrial luminal epithelium undergoes hyperproliferation, stromal cell compaction, decidualization, and increased vascularization [15–17]. These morphological changes are also associated with extensive changes in expression of genes having roles in immune modulation, proteolysis, transcription regulation, and cell signaling [15]. Furthermore, the expression of these genes is altered in the endometria of infertile women or those experiencing recurrent implantation failure [8, 18]. This has led to the emergence of a new step in our understanding of embryo implantation, where the receptive endometrium is further modulated by a blastocyst-driven signature to promote implantation [19]. However, limited information is available regarding the molecular features of this embryo-endometrium crosstalk.

The luminal epithelium of the endometrium is highly enriched in glycoproteins and have diverse functions [20]. Among these, Glycodelin and MUC1 are the most extensively studied glycoproteins for their roles in embryo implantation [20, 21]. Glycodelin-A is found abundantly in the secretory endometrium, decidua, and amniotic fluid that plays important roles in fertilization, embryo implantation, and placentation [22]. Its levels are altered in the endometria of women with endometriosis and recurrent implantation failure [23, 24]. MUC1 is abundantly expressed in the receptive-stage endometrium; its lower levels in serum or uterine flushing are associated with recurrent implantation failure [23, 25]. The other glycoproteins in the endometrial epithelium are the selectin and galectins that play role in embryo apposition and implantation [20, 25, 26]. Oviductal glycoprotein 1 (OVGP1) is a high-molecular-weight glycoprotein thought to be exclusively secreted by the non-ciliated epithelial cells of the fallopian tube [27, 28]. Identified first as an estrogen-induced secretory protein in the oviduct, OVGP1 aids in sperm capacitation, fertilization, and early embryonic development [29, 30]. The endometrial epithelium is a continuum of the oviductal epithelium and shares the same embryonic origin, and it is not surprising that several genes including steroid receptors, transcription factors, and growth factors are expressed commonly by both the uterine and oviductal epithelium [31]. In this context, OVGP1 is considered as an exception as its transcripts and protein are not detected in the endometrium of most species, except the hamster [29, 32, 33]. In our attempts to identify embryo-regulated molecules, we serendipitously discovered the expression of OVGP1 in the mouse endometrial epithelium during embryo implantation.

In the present study, we report a detailed analysis of the expression, regulation, and requirement of OVGP1 in the endometrial epithelial cells during embryo implantation. The

results reveal that in the endometrial epithelial cells, OVGP1 regulates the expression of receptivity-related genes and aids in trophoblast adhesion.

Materials and methods

Ethics approval

The study was approved by the Institutional Animal Ethics Committee (IAEC) of the National Institute for Research in Reproductive (NIRRH).

Animal/tissue collection

Ten-week-old C57BL/6 mice housed in an experimental animal facility of NIRRH were used. Female mice were mated with males of proven fertility, and the day of vaginal plug was defined as day 1. The mice were sacrificed on different days after detection of vaginal plug corresponding to various stages of embryo implantation [7]. The uteri were collected on day 4 morning (9:00 AM) when the embryo enters the endometrial lumen and it is in the receptive phase and day 4 evening (9:00 PM) when the embryo initiates apposition. The uteri were also collected on day 5 morning (9:00 AM) when the embryo attaches to the endometrial luminal epithelium and also day 5 evening (9:00 PM) when the embryo initiates epithelial breaching. The oviduct and uterus were also harvested from non-pregnant mice in both estrus and diestrus stages. Uteri from three mice were collected at each time points/stages as biological replicates point/stage. The tissues were fixed overnight in 4% paraformaldehyde (Sigma-Aldrich, MO, USA) at 4 °C and processed for paraffin embedding and sectioning.

Cell culture

Ishikawa cells are derived from well-differentiated adenocarcinoma of the endometrium. These cells express steroid receptors and mimic most features of the endometrial epithelium [34, 35]. Ishikawa cells were kindly provided to us by Dr. Geetanjali Sachdeva (NIRRH). The cells were cultured in DMEM/F12 with 10% fetal bovine serum (FBS) and 1% antibiotic-antimycotic (all from Gibco, Thermo Fisher Scientific, MA, USA). JAr cells are placental choriocarcinoma cell lines that were kindly provided to us by Dr. Geetanjali Sachdeva and were cultured in RPMI with 10% FBS and 1% antibiotic-antimycotic (all from Gibco). The cells were maintained in 5% CO₂ at 37 °C.

Hormonal treatment

Ishikawa cells were cultured in DMEM/F12 with steroid-stripped 2% FBS for 24 h and then treated with 17-estradiol

(10^{-8} M) and progesterone (10^{-6} M) (both from Sigma) alone or in combination as detailed previously [36]. Cells were treated with 1 IU of human chorionic gonadotropin (hCG) (Bharat Serum and Vaccines Ltd., Thane, Maharashtra, India) alone or in combination with estrogen and progesterone. Cells were harvested at 6, 12, 18, and 24 h of treatment. Cells without any treatment were harvested at each time point and were used as controls.

OVGP1 gene silencing

Gene silencing was performed using plasmid containing shRNA (pGIPZ vector) targeting *OVGP1* gene (V3LHS_361729; Dharmacon, Colorado, USA). As controls, a plasmid containing scrambled shRNA (V3LHS_65890S; Dharmacon) was used. Endotoxin-free plasmid was transfected in Ishikawa cells using Xtreme Gene HP transfection reagent (Sigma) as per the manufacturer's protocol. Post-72 h of transfection, the cells were treated 8 $\mu\text{g}/\text{ml}$ of puromycin (Gibco) for 15 days. The cells were then serially subcultured and harvested at 70% confluency for further experiments. The knockdown efficiency of *OVGP1* gene was examined by qPCR and immunofluorescence. All the experiments were performed between passages 30–35.

Total RNA isolation and real-time PCR

Chilled Trizol reagent (Invitrogen, Thermo Fisher Scientific) was directly added to the cells, and RNA was extracted as described previously [36]. RNA was treated with DNase I (New England Biolabs, MA, USA) and reverse-transcribed using high-capacity cDNA reverse transcriptase kit (Applied Biosystems, Thermo Fisher Scientific). Quantitative PCR was performed (in duplicates) in the CFX-96 thermal cycler (Bio-Rad, CA, USA) using iQ SYBR green chemistry (Bio-Rad). The amplification conditions for all genes (*GAPDH*, *HOXA10*, *HOXA11*, *IL6*, *IL15*, *IL11*, *ITGAV*, *ITGA5*, *ITGA6*, *ITGB3*, *LIF*, *MMP2*, *MMP9*, *OVGP1*, *PAEP*, *TGFB1*, *TIMP1*, *TIMP2*, *TIMP3*) were initial denaturation 95 °C for 30 s, specific annealing temperature (Supplementary Table 1) for 30 s, and extension at 72 °C for 30 s for 35 cycles. The fluorescence emitted at each cycle was collected for the entire period of 30 s during the extension step of each cycle. The homogeneity of the PCR amplicons was verified using the melt curve method. The sequences of the primers and annealing temperature used are given in Supplementary Table 1. Gene expression was normalized to *GAPDH*, and fold change was estimated using Livak's method [36].

Immunofluorescence

Five-micrometer-thick paraffin embedded sections were prepared on poly-L-lysine (Sigma)-coated slides (HiMedia, Maharashtra, India), deparaffinized in xylene, and rehydrated

in decreasing concentrations of alcohol. Immunofluorescence was performed as described previously [37]. In brief, the antigenic sites were unmasked by incubation of sections at 90 °C in Tris EDTA buffer (10 mM, pH 9) followed by incubation in 10 mM sodium borohydrate (Sigma) to reduce autofluorescence. The sections were blocked in 5% bovine serum albumin (BSA; MP Biomedicals, Maharashtra, India), followed by overnight incubation with the anti-OVGP1 (ab74544) antibody (Abcam, Cambridge, UK), at a dilution of 1:500 in phosphate-buffered saline (PBS). The sections were washed with PBS, and the bound antibodies were detected with Alexa Fluor 568 donkey anti-rabbit IgG (Molecular Probes, Thermo Fisher Scientific). The sections were counterstained in 0.1 $\mu\text{g}/\text{ml}$ of DAPI (Sigma) and mounted in the antifade medium. Imaging was done on inverted fluorescence microscope equipped with sCMOS camera (Leica Microsystems DMi8, Mannheim, Germany). The images were analyzed, and the quantification was done using LAS X software (Leica Microsystems).

Ishikawa cells were grown on a coverslip and fixed in absolute methanol for 20 min, followed by blocking in 5% BSA, and OVGP1 immunofluorescence was carried out as mentioned above.

Western blotting

Total cell lysates were prepared in ice-cold lysis buffer (150 mM NaCl, 50 mM Hepes, 5 $\mu\text{g}/\text{ml}$ digitonin, 0.1% Triton X-100) containing protease inhibitor (all from Sigma). Thirty-microgram protein was separated on 8% SDS gel, transferred overnight at 30 V onto a nitrocellulose membrane (GE Healthcare Life Sciences, IL, USA) in a transfer buffer (0.03% SDS and 10% methanol). The membrane was blocked in 5% non-fat dried milk (Nestle, Uttar Pradesh, India) and probed with anti-OVGP1 (1:500) antibody (Abcam). After extensive washings, the blots were probed with a secondary antibody labeled with horseradish peroxidase-conjugated anti-rabbit IgG (ABclonal, MA, USA). Detection was done using chemiluminescence kit (Pierce, Thermo Fisher Scientific).

Trophoblast adhesion assay

JAr cells were treated with conditioned media derived from *OVGP1* knockdown cells and scrambled cells for 24 h. JAr cells were trypsinized and allowed to adhere to a 96-well plate. At the end of 6 h, the wells were washed three times with PBS, and cells were fixed and stained with 0.5% crystal violet (Sigma). After extensive washing, the cells were lysed, and the absorbance was measured at 590 nm. The experiment was done using two independent pools of media derived from scrambled and *OVGP1* knockdown cells. Each experiment was carried out in triplicates.

Statistical analysis

The mean \pm SD for all the experimental data was calculated, and statistical analysis was done using GraphPad Prism, version 5, either by Student's *t* test or by one-way ANOVA using Tukey's all column comparison test. $p < 0.05$ was accepted as statistically significant.

Results

OVGP1 is expressed in the mouse endometrium at the time of implantation

By immunofluorescence, the signals for OVGP1 were detected exclusively in the luminal epithelial cells of the endometrium from the pregnant mice (Fig. 1). Faint OVGP1 staining was observed in the luminal epithelium on the morning of day 4, the intensity increased on the evening

of day 4 (Fig. 1a, b). This increase was not statistically significant (Fig. 1i). On the morning of day 5, OVGP1 expression increased further; and this increase was statistically significant (Fig. 1c, i). On the evening of day 5, the expression of OVGP1 appeared weaker as compared to day 5 morning; this reduction was statistically significant (Fig. 1d, i). In the oviduct (positive control), intense cytoplasmic staining for OVGP1 staining was detected; no staining was observed in the endometrium at both estrus and diestrus stages (Fig. 1e–g). No staining was observed in negative control without primary antibody (Fig. 1h).

OVGP1 is expressed in Ishikawa cells

A single band of expected size (172 bp) for *OVGP1* was detected by RT-PCR in cDNA of Ishikawa; no bands were observed in the reactions without reverse transcriptase. *GAPDH* is an internal control (Fig. 2a). The PCR products of *OVGP1* showed 100% sequence similarity to human *OVGP1* (data not shown). By

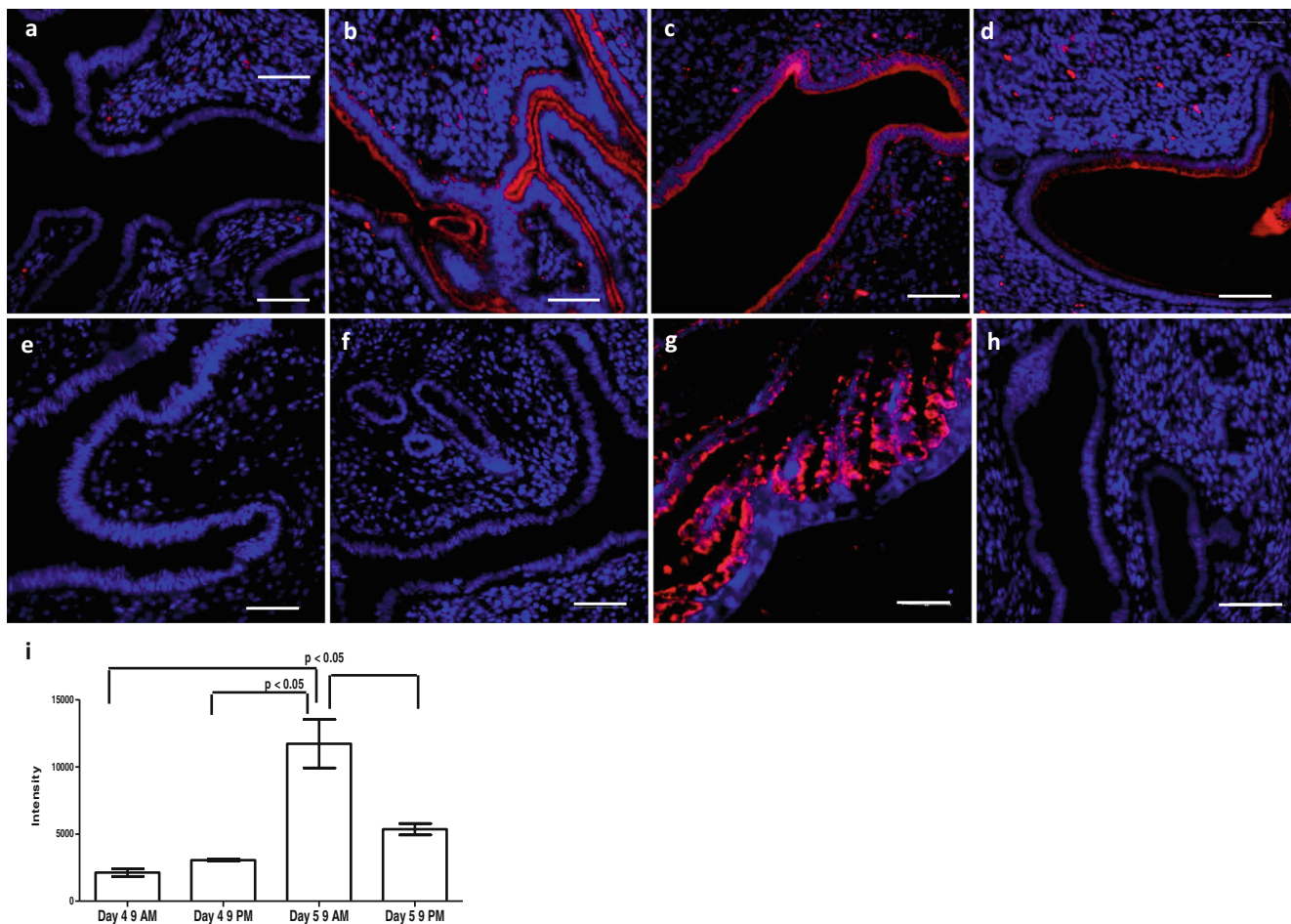


Fig. 1 Expression of OVGP1 in mouse endometrium at time of embryo implantation. Paraffin sections were probed with OVGP1 antibody and detected with Alexa Fluor 568. **a** Day 4 morning. **b** Day 4 evening. **c** Day 5 morning. **d** Day 5 evening. **e** Estrus stage. **f** Diestrus stage. **g** Oviduct. **h** Negative control (without primary antibody). Red staining is for OVGP1,

and blue is for the nuclei. Scale bar is 50 μ m. **i** Intensity of OVGP1 staining during the course of embryo implantation. Y-axis is the intensity values for OVGP1 staining. Data is mean \pm SD for the three biological replicates at each point. * indicates significant difference as compared to control ($p < 0.05$)

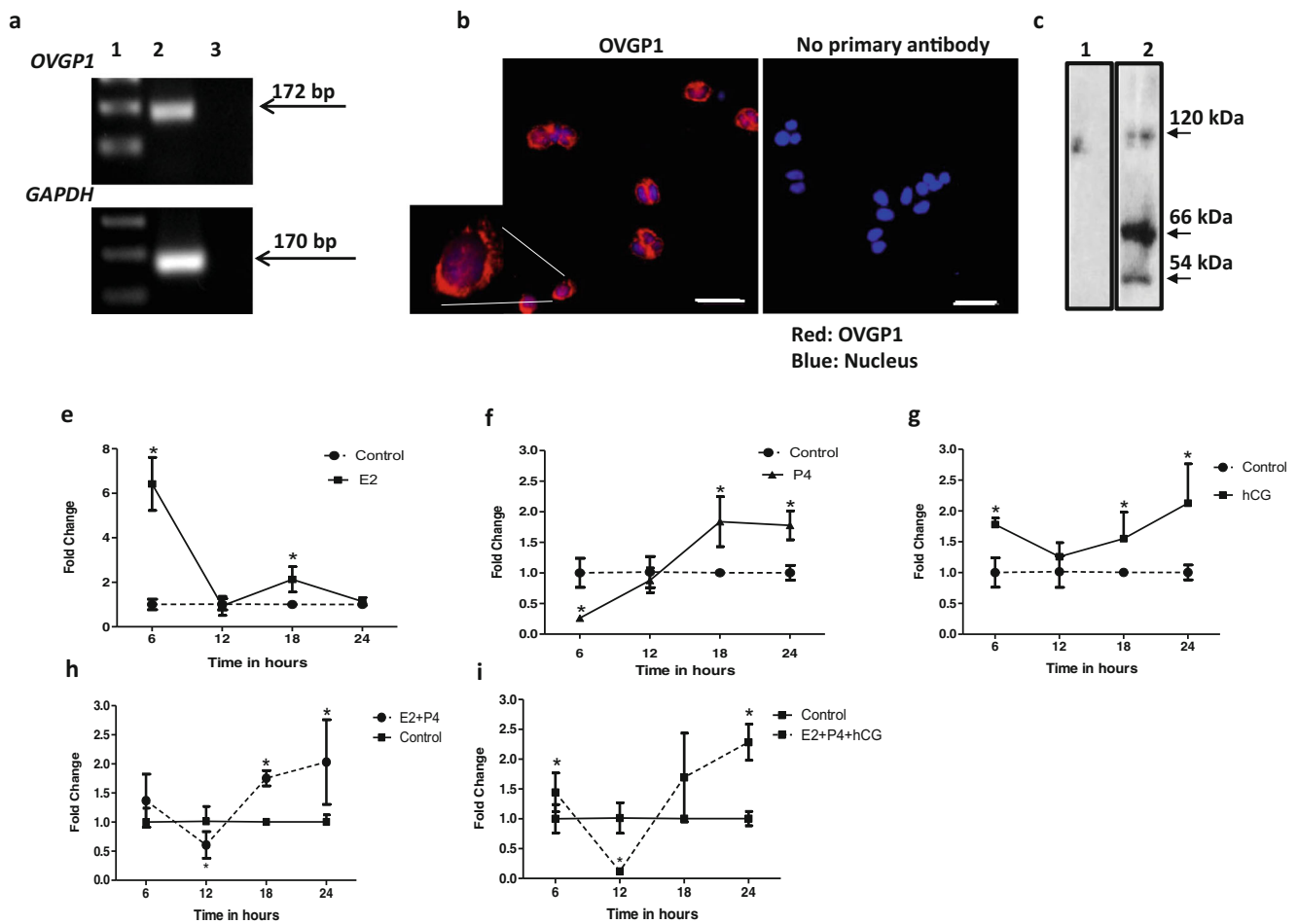


Fig. 2 Effects of estrogen, progesterone, and human chorionic gonadotropin (hCG) on *OVGP1* expression in human endometrial epithelial cells. **a** Expression of *OVGP1* mRNA from Ishikawa cells. Lane 1 is 100 bp ladder, lane 2 cDNA from Ishikawa mRNA, and lane 3 no reverse transcriptase control. The bands for *OVGP1* (172 bp) and *GAPDH* (170 bp) are shown by arrow marks. **b** Immunofluorescence for *OVGP1* in Ishikawa cells. Red staining is for *OVGP1*, and blue is for the nuclei. Scale bar is 50 μ m. A cell is digitally zoomed \times 5 times and shown. **c** Western blotting for *OVGP1* in protein isolated from Ishikawa cells. Lane 1 is negative control (incubated without primary antibody),

and lane 2 is sample incubated with primary antibody. The approximate size of the bands is marked with an arrow. All the experiments were done at least three times. Ishikawa cells were treated with estrogen (E₂), progesterone (P₄), and hCG alone or in combinations (e–i) for varying time points, and level of *OVGP1* mRNA was measured by real-time PCR. Y-axis is fold change as compared to untreated control at each time point (control value taken as 1). X-axes are time points in hours post-treatment. Values are mean \pm SD for six replicates. * indicates mean value significant different ($p < 0.05$) as compared to respective control

immunofluorescence, intense cytoplasmic staining for *OVGP1* was detected in Ishikawa cells; at higher magnification, the localization appears perinuclear (Fig. 2b). No *OVGP1* staining was observed in the cell incubated without primary antibody. By Western blotting, three bands of approximately 110–130 kDa, 60–66 kDa, and 54 kDa were detected in Ishikawa cell lysate probed with anti-*OVGP1* antibody. No bands were detected in negative control without primary antibody (Fig. 2c).

Steroid hormones and human chorionic gonadotropin (hCG) regulate expression of *OVGP1* in Ishikawa cells

As compared to controls, *OVGP1* mRNA levels increased significantly ($p < 0.05$) at 6 and 18 h of estrogen treatment; the levels were not significantly different at 12 and 24 h

(Fig. 2e). As compared to untreated controls, treatment with progesterone significantly reduced ($p < 0.05$) *OVGP1* levels at 6 h of treatment, which further increases significantly ($p < 0.05$) at 18 h and remained high at 24 h post-treatment (Fig. 2f). The levels of *OVGP1* were significantly ($p < 0.05$) high through the course of hCG treatment as compared to control, except for 12 h where the increase in expression was not significant (Fig. 2g).

As compared to controls, a combination of estrogen and progesterone significantly increased ($p < 0.05$) *OVGP1* levels at 18 and 24 h of treatment (Fig. 2h). As compared to controls, the inclusion of hCG along with estrogen and progesterone increased *OVGP1* mRNA levels at 6, 18, and 24 h. However, the increase was statistically significant ($p < 0.05$) only at 6 and 24 h (Fig. 2i).

At 12 h, the levels of *OVGP1* were significantly low ($p < 0.05$) as compared to controls.

Downregulation of *OVGP1* alters the expression of endometrial receptivity-related genes in Ishikawa cells

As compared to scrambled shRNA-transfected Ishikawa cells (controls), there was 60% reduction in *OVGP1* mRNA in cells transfected with shRNA against *OVGP1* (Fig. 3a). The intensity of OVGP1 protein was also reduced. There was 50% reduction in levels of OVGP1 protein in knockdown cells as compared to control (Fig. 3b, c). In both cases, the reduction was statistically significant ($p < 0.05$).

As compared to controls, the expression of *ITGAV*, *ITGB3*, and *ITGA5* was significantly ($p < 0.05$) reduced in cells knockdown for *OVGP1*; *ITGA6* mRNA levels were not significantly altered (Fig. 3d–g). The expression of

HOXA10 mRNA was reduced significantly ($p < 0.05$), whereas *HOXA11* mRNA levels were significantly ($p < 0.05$) increased in *OVGP1* knockdown cells as compared to controls (Fig. 3h, i).

Downregulation of *OVGP1* alters expression of cytokines, matrix metalloproteases, and tissue inhibitor of matrix metalloproteases in Ishikawa cells

As compared to controls, the mRNA levels of *LIF* and *IL15* were significantly ($p < 0.05$) lowered in Ishikawa cells knockdown for *OVGP1* (Fig. 4a, d). The expression of *IL6*, *IL11*, *TGFB1*, and *PAEP* was not significantly altered in *OVGP1* knockdown cells as compared to controls (Fig. 4b–f). As compared to control, there was significant ($p < 0.05$) increase in the levels of *MMP9*, *TIMP1*, and *TIMP3* in *OVGP1* knockdown cells; the levels of *MMP3* and *TIMP2* were unaltered (Fig. 5).

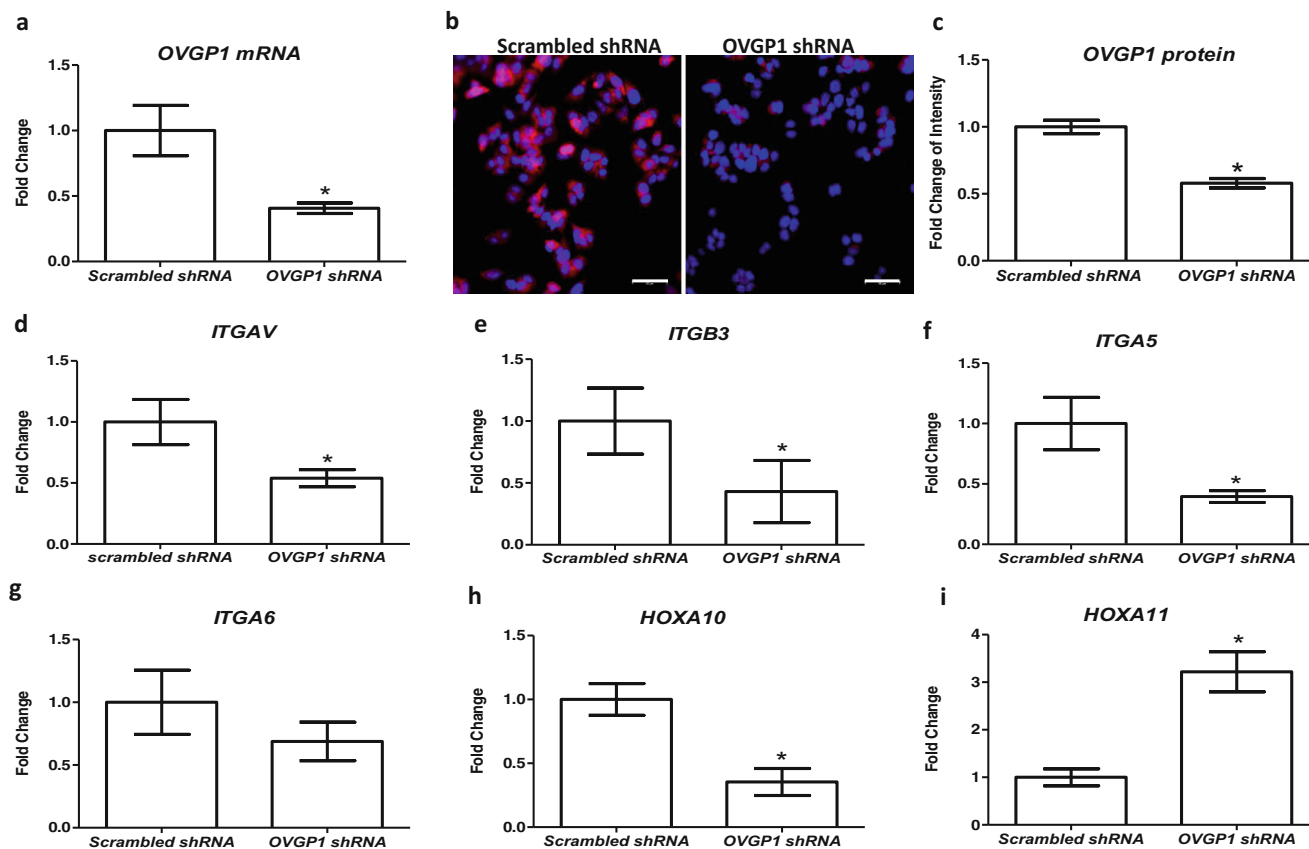


Fig. 3 Effect of loss of *OVGP1* on expression of receptivity markers in the endometrial epithelial cells. **a** mRNA levels of *OVGP1* in scrambled and *OVGP1* shRNA-transfected Ishikawa cells. Y-axis is fold change where values obtained from scrambled cells were taken as 1. **b** Immunofluorescence for *OVGP1* in scrambled and knockdown Ishikawa cells. Red staining is for *OVGP1*, and blue is for the nuclei. Scale bar is 50 μm . **c** Intensity of *OVGP1* staining in scrambled and *OVGP1* shRNA-transfected Ishikawa cells. Y-axis is intensity values for

OVGP1 staining. In both **a** and **b**, data is mean \pm SD for the three independent replicates. * indicates significant difference as compared to control ($p < 0.05$). Effect of loss of *OVGP1* on mRNA levels of *ITGAV*, *ITGB3*, *ITGA5*, *ITGA6*, *HOXA10*, and *HOXA11* (**d**–**i**). Y-axis is fold change where values obtained from scrambled cells were taken as 1. Data is the mean \pm SD for the three independent replicates. * indicates significant difference as compared to control ($p < 0.05$)

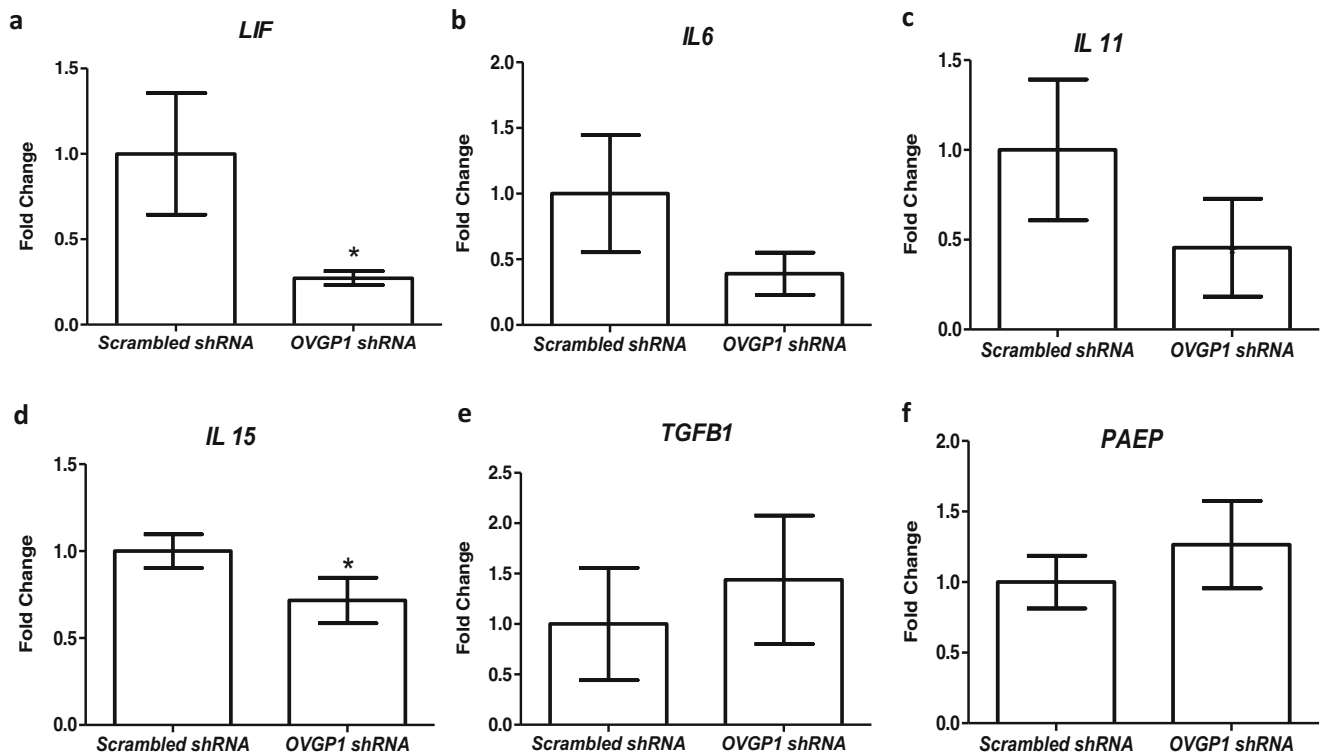


Fig. 4 Effect of loss of *OVGP1* on mRNA levels of *LIF*, *IL6*, *IL11*, *IL15*, *TGFB1*, and *PAEP* (a–f). Y-axis is fold change where values obtained from scrambled cells were taken as 1. Data is the mean ± SD for the three independent replicates. * indicates significant difference as compared to control ($p < 0.05$)

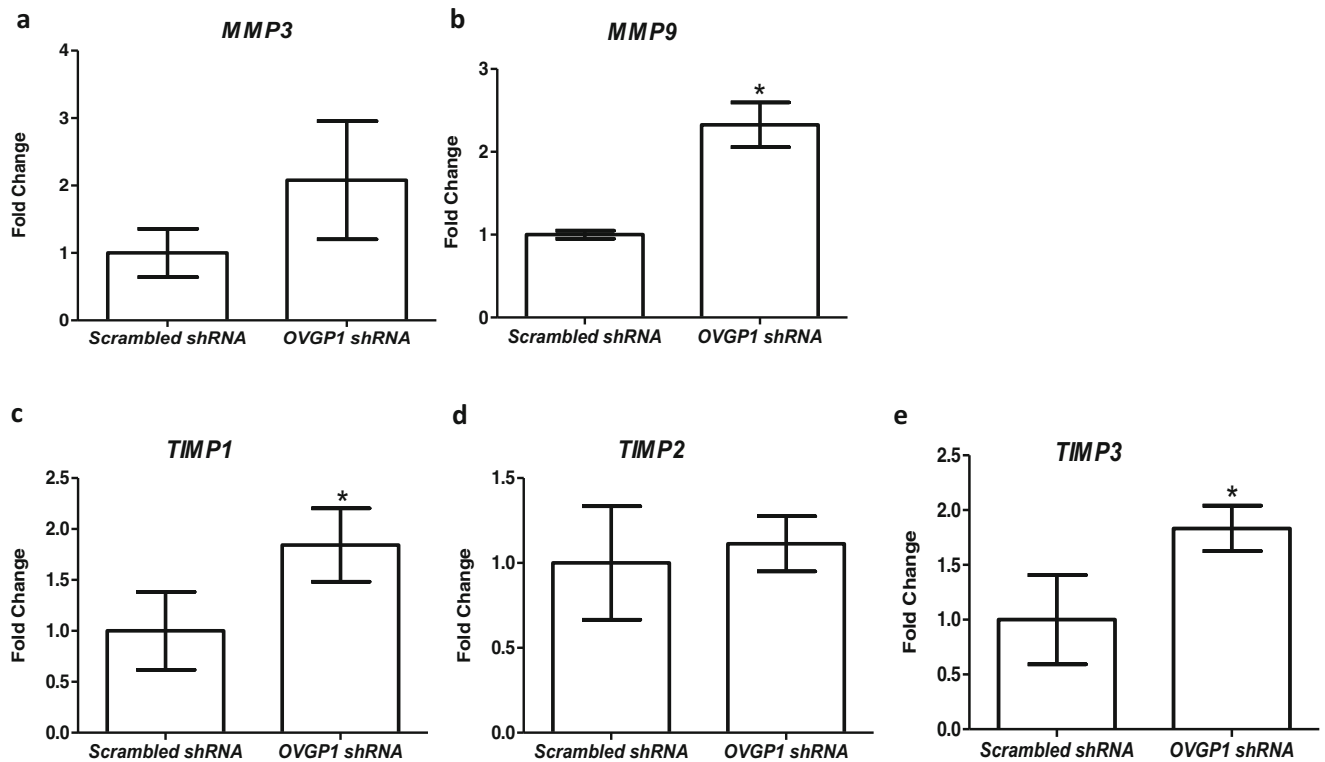


Fig. 5 Effect of loss of *OVGP1* on mRNA levels of *MMP3*, *MMP9*, *TIMP1*, *TIMP2*, and *TIMP3* (a–e). Y-axis is fold change where values obtained from scrambled cells were taken as 1. Data is the mean ± SD for the three independent replicates. * indicates significant difference as compared to control ($p < 0.05$)

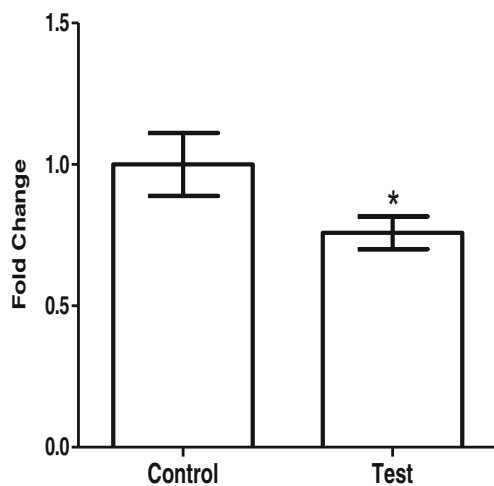


Fig. 6 Effect of loss of *OVGP1* in the endometrial epithelial cells on adhesion of trophoblast cells. Control is conditioned media from Ishikawa cells transfected with scrambled shRNA. Test is conditioned media from Ishikawa cells transfected with *OVGP1* shRNA. Trophoblast cells (JAR cell line) were treated with the conditioned media for 24 h, trypsinized, and allowed to adhere for 6 h. The number of adherent cells was measured as described in the “Materials and methods” section. Values on Y indicate fold change where values obtained from control were taken as 1. Data are the mean \pm SD of absorbance from two independent experiments done in triplicates. * indicates significant difference as compared to control ($p < 0.05$)

Loss of *OVGP1* in Ishikawa cells reduces the adhesion of trophoblast cells

As compared to JAR cells incubated with conditioned media from scrambled shRNA-transfected Ishikawa cells, there was a decrease in adhesion of JAR cells treated with conditioned medium from *OVGP1* knockdown cells (Fig. 6). There was 30% reduction in the adhesion, and this was statistically significant ($p < 0.05$).

Discussion

The results of the present study demonstrate that *OVGP1* is expressed exclusively in the luminal epithelial cells of the mouse at the time of embryo apposition. Further, we demonstrate that loss of *OVGP1* in human endometrial epithelial cells alters the expression of genes involved in endometrial receptivity. Finally, the results also reveal that culture medium derived from epithelial cells knockdown for *OVGP1* reduces the in vitro adhesiveness of trophoblast cells.

Previous studies have demonstrated that *OVGP1* is not expressed by the cycling endometrium of mice and monkeys [27, 29, 32]. However, microarray analysis of isolated cells from human endometrium reveals significant enrichment of *OVGP1* mRNA in the epithelial cells (Supplementary Fig. 1). Interestingly, in the present study, we observed that in the presence of the embryo, *OVGP1* protein is expressed by the

luminal epithelial cells at the time of implantation. This gain of expression is due to the transcription of *Ovgp1* by the endometrial cells as its mRNA is detected in the endometria of pregnant mice (Supplementary Fig. 2). A similar increase in *OVGP1* expression is observed in the luminal epithelial cells at the time of embryo apposition in the hamster [33]. These results together suggest that *OVGP1* is induced in the endometrial epithelium at the time of implantation.

Since *OVGP1* is induced by the endometrial epithelial cells in the presence of the embryo, we next asked if embryonic signals induce *OVGP1* expression in the endometrial epithelium. For this purpose, we screened several cell lines of human endometrium and observed that Ishikawa cells express significant amounts of *OVGP1* mRNA and protein; three bands of *OVGP1* corresponding to pre-processed (60–66 kDa), processed (54 kDa), and glycosylated (110–130 kDa) forms were detected. Furthermore, the protein was localized in the perinuclear region; *OVGP1* was also detected on the surface of Ishikawa cells and conditioned medium (data not shown) suggesting that *OVGP1* is transcribed, translated, and secreted by Ishikawa cells. Thus, Ishikawa cells were chosen to carry out further studies on regulation and roles of *OVGP1*. Embryo implantation involves coordinated signaling from the embryo and also the ovarian steroids estrogen and progesterone. It is known, that in the progesterone-primed endometrium, a peak in estrogen is essential to promote implantation, and this peak of estrogen brings about the coordinated expression of several genes essential for receptivity [3–7, 38]. Furthermore, this steroid-driven signature of the endometrium is modulated by hCG secreted from the embryo [15–17]. Thus, we studied the effects of estrogen, progesterone, and hCG alone or in combination to test the regulation of *OVGP1* in the endometrial epithelium. Treatment of estrogen transiently increases the *OVGP1* expression in Ishikawa cells. Interestingly, in the human endometrium, *OVGP1* mRNA levels are highest in the proliferative phase which coincides with estrogen peak (Supplementary Fig. 3). Indeed, in the oviduct, *OVGP1* is known to be induced by estrogen [29], and *OVGP1* promoter contains ten half-palindromic estrogen responsive elements [28]. Interestingly, along with estrogen, hCG consistently and significantly increased the *OVGP1* expression in Ishikawa cells. These results suggest that the increase in *OVGP1* observed in the luminal epithelium at the time of embryo implantation is perhaps due to the combined effect of estrogen and hCG.

We next assessed the functional significance of *OVGP1* in the endometrial epithelial cells in the context of embryo implantation. For this purpose, endogenous expression *OVGP1* was silenced in Ishikawa cells using shRNA, and the expression of genes associated with receptivity was studied. Integrins are required for embryo implantation, and the aberrant expression of integrins is observed in the endometria of women with unexplained infertility [39, 40]. *HOXA10* and *HOXA11* are essential for uterine receptivity and mice null

for both these are infertile due to the failure of the embryo to implant (reviewed in [41, 42]). In addition, women with unexplained infertility have reduced expression of *HOXA10* in their uteri [42]. From our study, it appears that *OVGP1* is required to maintain the expression of integrins and *HOXA10* in the endometrial epithelium, as inhibition of *OVGP1* expression reduced the mRNA levels of *HOXA10*, *ITGAV*, *ITGA5*, and *ITGB3*. Interestingly, at the same time, the levels of *HOXA11* were elevated upon the reduction of *OVGP1*; the levels of *PAEP* and *TGFBI* were unaltered. Since *HOXA10* controls integrin expression and permits pinopode formation [43, 44], it is possible that *OVGP1* aids this process by maintaining the levels of these molecules in the receptive endometrium. Together, these results suggest that *OVGP1* is required to maintain the balance of receptivity-related molecules in the endometrial epithelium.

Cytokines play an integral role in embryo implantation and stromal cell decidualization. *LIF*, *IL6*, *IL11*, and *IL15* are secreted by the uterine epithelium that are required for decidualization; they also promote trophoblast proliferation, adhesion, and invasion [45–50]. Thus, we tested if *OVGP1* has any role in the regulation of cytokine expression. The results revealed that the levels of *LIF* and *IL15* were reduced in cell knockdown for *OVGP1*; the levels of *IL6* and *IL11* were also low, albeit not significantly. These results indicate that endometrial *OVGP1* might aid embryo implantation by controlling cytokine expression for decidualization and trophoblast invasion.

One of the requirements for embryo implantation is degradation of extracellular matrix proteins of the endometrium. This is achieved by increasing the production of proteases such as matrix metalloproteases (MMPs) and decreasing the activity of tissue inhibitors of matrix metalloproteases (TIMPs). To test if *OVGP1* has any role in the regulation of protease activity at the time of implantation, we measured the mRNA levels of *MMP3*, *MMP9*, *TIMP1*, *TIMP2*, and *TIMP3* in Ishikawa cells knockdown for *OVGP1*. The results revealed an increase in the expression of *MMP9*, *TIMP1*, and *TIMP3* in response to the loss of *OVGP1*. *MMP9* is a key enzyme required for degradation of collagenase, and an increase in its expression is associated with loss of tissue integrity [51]. Thus, an excess of *MMP9* might be detrimental for embryo and endometrial cells. While TIMPs are considered as an antagonist for MMPs, mice transgenic for *TIMP1* have smaller deciduas and higher resorption rates [52]. These observations suggest that high *TIMP1* is also detrimental for implantation. Excess of *TIMP3* is known to inhibit matrix degradation at the time of implantation and trophoblast invasion [46]. Thus, it is plausible that *OVGP1* in the epithelium may be required to balance the production of MMPs and TIMPs and control ECM remodeling for implantation.

For successful implantation, the embryonic trophoblast must adhere to the endometrial epithelium. While the role of

epithelial integrins and membrane-bound proteins in the process of trophoblast adhesion is undisputable [6, 21], the process is also governed by secretory growth factors, cytokines, chemokines, and other molecules [37, 47, 50]. From our results, it is evident that the levels of cytokines, MMPs, and TIMPs are altered in epithelial cells knockdown for *OVGP1*. Since these molecules are known to alter trophoblast physiology [46], we asked if secretions from *OVGP1* knockdown cells have any effects on trophoblast adhesion. Indeed, trophoblast cells treated with conditioned medium of *OVGP1* knockdown cells had lower adhesiveness suggesting that epithelial *OVGP1* probably maintains a secretome that is conducive for embryonic attachment. However, it is intriguing to note that mouse knockout for *Ovgp1* is fertile, but they have reduced litter size [53]. Also, *OVGP1* transcripts are significantly reduced in the secretory phase endometria of women with recurrent implantation failure (Supplementary Fig. 4). Based on these results, we suggest that *OVGP1* might have critical roles in embryo implantation.

To summarize, in the present study, for the first time, we demonstrate that *OVGP1* is specifically induced in the luminal epithelium at the time of embryo implantation; this increase in *OVGP1* is required to maintain the balanced expression of receptivity-related genes and aids in trophoblast adhesion. This study adds one more molecule to the growing list of embryo-induced factors in endometrium required for implantation. We believe that identification of such factors will expand our knowledge of the mechanisms involved in embryo implantation, and in long-term such studies might aid in improving pregnancy rates of assisted reproduction.

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Compliance with ethical standards

The study was approved by the Institutional Animal Ethics Committee (IAEC) of the National Institute for Research in Reproductive (NIRRH).

Conflict of interest The authors declare that they have no conflict of interests.

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