

# Proliferative potential and phenotypic analysis of long-term cultivated human granulosa cells initiated by addition of follicular fluid

Lenka Bruckova · Tomas Soukup · Benjamin Visek · Jiri Moos · Martina Moosova ·  
Jana Pavelkova · Karel Rezabek · Lenka Kucerova · Stanislav Micuda ·  
Eva Brackova · Jaroslav Mokry

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## Abstract

**Purpose** The aim of this study was to develop and optimize a strategy for long-term cultivation of luteinizing human granulosa cells (GCs).

**Methods** GCs were cultivated in DMEM/F12 medium supplemented with 2% fetal calf serum. In vitro prolifera-

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**Capsule** This paper is especially focused on the improvement of the cultivation protocol enabling granulosa cells to proliferate and maintain the typical phenotypic expression.

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L. Bruckova  
Department of Analytical Chemistry, Faculty of Chemical  
Technology, University of Pardubice,  
Pardubice, Czech Republic

L. Bruckova · T. Soukup (✉) · B. Visek · J. Mokry  
Department of Histology and Embryology,  
Charles University in Prague, Medical Faculty in Hradec Kralove,  
Simkova 870,  
500 38 Hradec Kralove, Czech Republic  
e-mail: soukupto@lfhk.cuni.cz

J. Moos · M. Moosova · J. Pavelkova · K. Rezabek  
Department of Obstetrics and Gynecology,  
Charles University in Prague,  
1st Medical Faculty and General Teaching Hospital in Prague,  
Prague, Czech Republic

L. Kucerova  
Department of Clinical Genetics,  
Teaching Hospital in Hradec Kralove,  
Hradec Kralove, Czech Republic

S. Micuda · E. Brackova  
Department of Pharmacology, Charles University in Prague,  
Medical Faculty in Hradec Kralove,  
Prague, Czech Republic

tion of GCs was supported by follicular fluid as well as FSH and growth factors.

**Results** The cultured GCs were maintained for 45 days with a doubling time of  $159 \pm 24$  h. GCs initiated by the addition of follicular fluid and cultivated under low serum conditions reached  $10 \pm 0.7$  population doublings. GCs maintain the typical phenotypic expression and the telomere length according to specific culture conditions.

**Conclusion** Our present study has demonstrated that GCs can be maintained in vitro for at least 45 days and this cell model can be beneficial when studying hormonal regulation associated with follicular maturation and preparation of oocytes for fertilization.

**Keywords** Cultivation · Follicular fluid · Granulosa cells · Phenotype

## Abbreviations

bFGF	basic fibroblast growth factor
COV434	cell line derived from a human granulosa cell tumor (Sigma-Aldrich)
DMEM/F12	dulbecco's modified eagle's medium
DT	doubling time
EGF	epidermal growth factor
FCS	fetal calf serum
FF	follicular fluid
FSH	follicle stimulating hormone
GCs	luteinizing human granulosa cells
GC8	one of the isolated GCs from patient with different characters
IVF	in vitro fertilization
LH	luteinizing hormone
PD	the number of population doublings

## Introduction

Human luteinizing granulosa cells (GCs) with follicular fluid (FF) are usually harvested during in vitro fertilization (IVF) from the preovulatory follicles of hormonally stimulated women. These cells are challenged with a peak of luteinizing hormone or human chorionic gonadotrophin, which may be regarded as a signal to convert GCs to luteal cells (1, 2). Each primordial follicle contains an oocyte, arrested in meiosis and surrounded by a single layer of flattened epithelial pregranulosa cells (3). The GCs continue to divide, and as follicle maturation is reached just before ovulation, GCs division decreases and the next phase of the oocyte meiosis resumes (4). Luteinized GCs are considered to be terminally differentiated, being replaced in the midluteal phase of the menstrual cycle by small, luteinized cells originating from the surrounding theca (5).

Wen et al. examined FF and GCs production of steroid from IVF patients. Despite the size of follicle triggered after controlled luteinisation, the levels of progesterone and testosterone were maintained at relatively constant levels (median 98.1  $\mu\text{mol/L}$  for progesterone, and 5.8 nanomoles/L for testosterone). However, estradiol levels were slightly lower in the larger follicles (follicular diameter 10–15 mm,

median 25.3 nanomoles/L; follicles  $\geq 15$  mm, median 15.1 nanomoles/L). Absolute steroid levels are associated with follicular size, not oocyte maturation/ability to fertilize (6). Estradiol, progesterone and testosterone are the main steroid hormones that play essential roles during the follicular and luteal phases of the menstrual cycle. However other cytokines, such as inhibins, activins, insulin growth factor-2, insulin growth factor binding proteins, tumour growth factor- $\beta$  and endothelial growth factor, have been measured in FF and correlated with oocyte maturation (6–8). The cellular content of FF aspirated during oocyte collection for assisted reproduction consisted of a mixture of GCs, erythrocytes and large vaginal epithelial cells (9, 10). Samples of FF used in this study were obtained from ovarian follicles of women undergoing IVF cycle in assisted reproduction centre. Only the samples of FF from cycles with mature and morphologically normal oocytes were included. Moreover, oocytes obtained in those cycles underwent IVF and/or ICSI and in all of the cycles, oocytes were fertilized and developing embryos were obtained (Table 1).

Disruptions of hormonal stimulation and communication between GCs and oocyte represent a serious complication of assisted reproduction. An example of such complications may be the development of ovarian hyperstimulation

**Table 1** Summary of antibodies used for flow cytometry to their specifications. BioLegend (BL), Beckman Coulter (BC), Caltag (C), BD Pharming (BD), eBioscience (eB), Sigma-Aldrich (SA). Mesen-

chymal marker (MM), endothelial marker (EM), haemopoietic marker (HM), adhesion molecule (AM), proliferation marker (PM)

	Clone	Function	
CD29	BL	TS2/16	MM, b1 integrin, associates with CD49a in VLA-1 integrin
CD31	BL	WM59	EM, PECAM-1, tissue regeneration and safely removing neutrophils
CD34	BC	581	HM, HPCA1, expression on early hematopoietic and vascular-associated tissue
CD44	C	MEM-85	MM, binds hyaluronic acid, mediates adhesion of stromal cells and ECM
CD45	C	HI30	HM, protein tyrosine phosphatase, receptor type C
CD49d	BC	HP2/1	AM, VLA-4, $\alpha 4$ integrin, associates with CD29, binds fibronectin, VCAM-1, adhesion
CD49e	BC	SAM-1	AM, VLA-5, $\alpha 5$ integrin, associates with CD29, binds fibronectin, adhesion, apoptosis
CD71	BL	MEM-75	PM, transferrin receptor 1 controls iron uptake during cell proliferation
CD73	BD	AD2	MM, ecto-5-nucleotidase, dephosphorylates nucleotides to allow nucleoside uptake
CD90	BC	F15-42-1-5	MM, Thy-1, possible inhibition of stem cell and neuron differentiation
CD105	BL	MEM-226	MM, EM, endoglin, protective role against pro-apoptotic factors
CD106	BL	STA	EM, VCAM-1, adhesion of cells to vascular endothelium
CD146	BC	TEA1/34	EM, MCAM, mesenchymal stem cells with greater differentiation potential
CD166	BC	3A6	MM, AM, ALCAM, ligand for CD6
CD184	C	12 G5	MM, HM, CXCR4, alpha-chemokine receptor specific for stromal-derived-factor-1
CD197	eB	3D12	Chemokine, CCR7
CD222	eB	MEM-238	PM, receptor for insulin-like growth factor 2
HLA-I	C	Tu149	Major histocompatibility class I
HLA-II	C	Tu36	Major histocompatibility class II
FSH-R	SA	F3929	PM, receptor FSH, GCs are the only cell type expressing the FSH-R
LH-R	SA	L6792	receptor LH, luteinization GCs

syndrome. No animal or human cellular model is available to study these specific health problems. GCs obtained during the collection of oocytes in the IVF cycle could represent a suitable in vitro model. Luteinized GCs are believed to be terminally differentiated, which significantly complicates the establishment and cultivation of GCs primary culture, and this is not supportive for long-term cultivation. The aim of our study was to find an optimal protocol for the long-term cultivation of GCs and to characterize the GCs phenotype in vitro.

## Material and methods

### Isolation of human granulosa cells

Human GCs were recovered from women undergoing IVF procedures after ovarian stimulation and ovulation induction. GCs with FF were obtained from 25 patients with their consent according to the guidelines of the Assisted Reproduction Center, Prague. Cultures of GCs were established using cells recovered from the FF during oocyte retrieval procedures. The mean age of patients was  $33.3 \pm 4$  year (range from 23–43 year), average BMI was  $24.7 \pm 4$ . Patients were stimulated by 10,000 IU Pregnyl (hCG, N.V. Organon Oss, Netherlands) 36 h prior to oocyte collection. An average of  $9 \pm 6$  ovarian follicles were harvested together with  $6 \pm 4$  oocytes from each patient. The mean value of FF was  $17 \pm 13$  ml. Samples that contained high quantities of erythrocytes were excluded. The GCs suspension was transported in FF (37°C), cells were centrifugated at 357 g for 5 min, then resuspended in cultivation medium and seeded on untreated plastic 12-well plates (TPP, Switzerland).

For GCs long-term cultivation, a new low-FCS cultivation medium was designed. Medium consisted of Dulbecco's Modified Eagle's Medium (DMEM/F12, Sigma-Aldrich, USA), 2% fetal calf serum FCS (PAA, Austria), 10 mg/ml ascorbic acid (Sigma-Aldrich, USA), 0.05  $\mu$ M dexamethasone (Sigma-Aldrich, USA), 200 mM L-glutamine (Invitrogen, USA), 10 mg/ml gentamycin (Invitrogen, USA), 10,000 units/ml penicillin and 10,000  $\mu$ g/ml streptomycin (Invitrogen, USA). Moreover we added 20 ng/ml EGF, 50 ng/ml bFGF (PeproTech, USA) and FSH (Puregon; NV Organon, Oss, the Netherlands).

On the day when the GCs were seeded, cultivation medium was supplemented with FF (2:1) and 2.5 mg/ml amphotericin (Sigma-Aldrich, USA). Cultivation medium was changed every day. Cells were cultivated at 37°C under aerobic conditions (5% CO<sub>2</sub>). Once adherent cells were more than 80% confluent, they were detached with 0.05% trypsin-EDTA (Invitrogen, USA) for 10 min and counted using a Z2 counter or cell viability analyser Vi-Cell XR 2.03 (both Beckman Coulter, USA).

### Immunofluorescence

For identification of cell surface antigens, cells were fixed with 4% paraformaldehyde at 20°C for 10 min. Blocking and diluent solution consisted of phosphate-buffer saline (PBS), 5% serum (Sigma) from the same species as was the primary antibody for 30 min. The samples were then incubated with a primary antibody for 90 min. Primary antibodies used in this study were specific for the following antigens: FSHR, LHR, nestin. After washing with PBS, the antigen-binding sites were visualized with a secondary antibody conjugated to fluorochromes. Goat anti-rabbit IgG coupled to Alexa 488 (1:300, Invitrogen) was used (incubation 30 min). Cells stained with omission of a primary antibody were used as negative controls. Cell nuclei were counterstained with DAPI (4'-6-diamidino-2-phenylindole, Sigma) for 5 min. Coverslips were mounted in Mowiol or polyvinyl/alcohol/glycerol with 1,4-diazobicyclo-[2.2.2]-octane (DABCO) as anti-fading agent. Samples were examined with a BX51 microscope (Olympus) equipped with a DP71 digital camera. The images were prepared using Adobe Photoshop software.

### Doubling time and population doublings

The number of population doublings was calculated by the following formula:

$$\text{No(PD)} = (\log N_t - \log N_0) / 0.301$$

No (PD) [number of population doublings], N<sub>t</sub> [number of cells after enzymatic dissociation/collection of cells], N<sub>0</sub> [number of cells seeded]. The resulting number of population doublings is assessed individually or cumulatively during cell cultivation. Regression analysis of individual values at every passage shows the trend of proliferative potential of cell culture. The cumulation of data is important to record Hayflick's limit (60 doubling of the population), i.e. to show an unlimited proliferative potential (11).

Time required to double the population is expressed in the equation:  $DT = (t \times \log 2) / (\log N_t - \log N_0)$ . DT [doubling time—time needed to double the population], t [elapsed time between seeding and cell harvesting], N<sub>t</sub> [number of cells after enzymatic dissociation/collection of cells], N<sub>0</sub> [number of cells seeded].

### Cultivation human granulosa tumor cells COV434

COV434 cells (Sigma-Aldrich, USA) were derived from a human granulosa cell tumor (12), but possess many characteristics of normal GCs (13). The line of immortalized granulosa cells (COV434) was established from a primary human granulosa cell tumor in 1984 from a 27 year

old female. The tumor granulosa cells were cultured in Dulbecco's modified Eagle's medium supplemented with 10% (v/v) FCS and 2 mM L-glutamine. Cultivation medium was changed every day.

### Karyotyping

Cells (subcultured in early passages and after reaching passage no. 5, 10, 15, 20 and 25) after 48 h of cultivation were subjected to a 4 h Demecolcemid (Sigma-Aldrich, USA) incubation followed by trypsin-EDTA detachment and lysis with hypotonic KCl and fixation in acid/alcohol. Metaphases were analyzed after GTG banding using the software Ikaros 5.0 (MetaSystems, USA).

### Quantitative PCR—telomere length measurement

Genomic DNA was extracted from the cells using the silica-gel-membrane-based DNeasy Tissue Kit (Qiagen, Melbourne, Australia). Telomere length measurement was performed by qPCR assay according to the method described by Cawthon (14) with small modifications. Briefly, the telomere repeat copy number to single-copy gene (T/S) ratio was determined using the equation:  $T/S = 2^{-\Delta Ct}$  (where  $\Delta Ct = Ct_{\text{telomere}} - Ct_{\text{single-copy gene}}$ ). Then, the T/S for each sample was normalized to T/S value of a reference DNA sample to standardize between different runs, i.e.  $-\Delta\Delta Ct$  was calculated for each sample. This value is proportional to the average telomere length of the evaluated sample. 36B4, encoding acidic ribosomal phosphoprotein P0 was used as the single copy gene. Telomere and 36B4 gene PCRs were always done in separate 96-well plates with each sample run in triplicate in the same well position on an ABI 7500 HT Detection System (Applied Biosystems, USA). Each 20  $\mu$ L reaction consisted of 20 ng DNA, 1 $\times$ SYBR Green master mix (Applied Biosystems), 200 nM telomere forward primer (CGG TTT GTT TGG GTT TGG GTT TGG GTT TGG GTT TGG GTT) and 200 nM telomere reverse primer (GGC TTG CCT TAC CCT TAC CCT TAC CCT TAC CCT). For the 36B4 gene PCR the following primer pairs were used: 36B4u, CAGCAAGTGGGAAGGTGTAATCC; 36B4d, CCCATTCTATCATCAACGGGTACAA. The DNA quantity standards were derived from serial dilutions of a reference DNA sample to produce three final concentrations (0.02, 0.2, and 2.0 ng/ $\mu$ L). In each run, a standard curve and a negative control (water) were included. Cycling conditions (for both telomere and 36B4 products) were 10 min at 95°C, followed by 45 cycles of 95°C for 15 s and 60°C for 1 min. Following amplification, a dissociation curve was done to confirm the specificity of the reaction. Standard and dissociation curves were generated with the ABI Prism 7500 SDS software.  $R^2$  for each standard curve was  $>0.98$ .

The Du145 prostate cancer cells were cultured as described previously (15) and served as a control for the comparison of the length of short telomeres with the relative T/S value of GCs.

### Flow cytometric analysis

A total of 100000 cells in 100  $\mu$ L PBS (Invitrogen, USA) per well of 96-well plate (TPP, Switzerland) were incubated with 1  $\mu$ g antibody for 30 min in the dark. Anti-human FSHR and LHR polyclonal antibodies (both Sigma-Aldrich, USA) were conjugated using a Lightning-Link<sup>TM</sup> Fluorescein Conjugation Kit (Innova Biosciences Ltd, UK) before use (according to the instructions of the manufacturer). Control samples included an autofluorescence control and an isotype (IgG1, IgG2a, IgG2b) immunoglobulin control conjugated with FITC and PE. Then all samples were analyzed using a flow cytometer (Cell Lab Quanta<sup>TM</sup> SC, Beckman Counter, USA). Data analysis was performed using CXP analytic software (Beckman Counter, USA). The percentage of positive cells was determined as a percentage of cells with a higher fluorescence intensity than the upper isotype immunoglobulin control of 0.5%. Classification criteria:  $<10\%$  no expression,  $10\text{--}40\%$  low expression,  $40\text{--}70\%$  moderate expression and  $>70\%$  high expression.

### Statistical analysis

The relationship between the average telomere length and the selected parameters was performed by correlation analysis using the statistic software GraphPad Prism 5.01 (San Diego, USA). A value of  $p < 0.05$  was considered statistically significant.

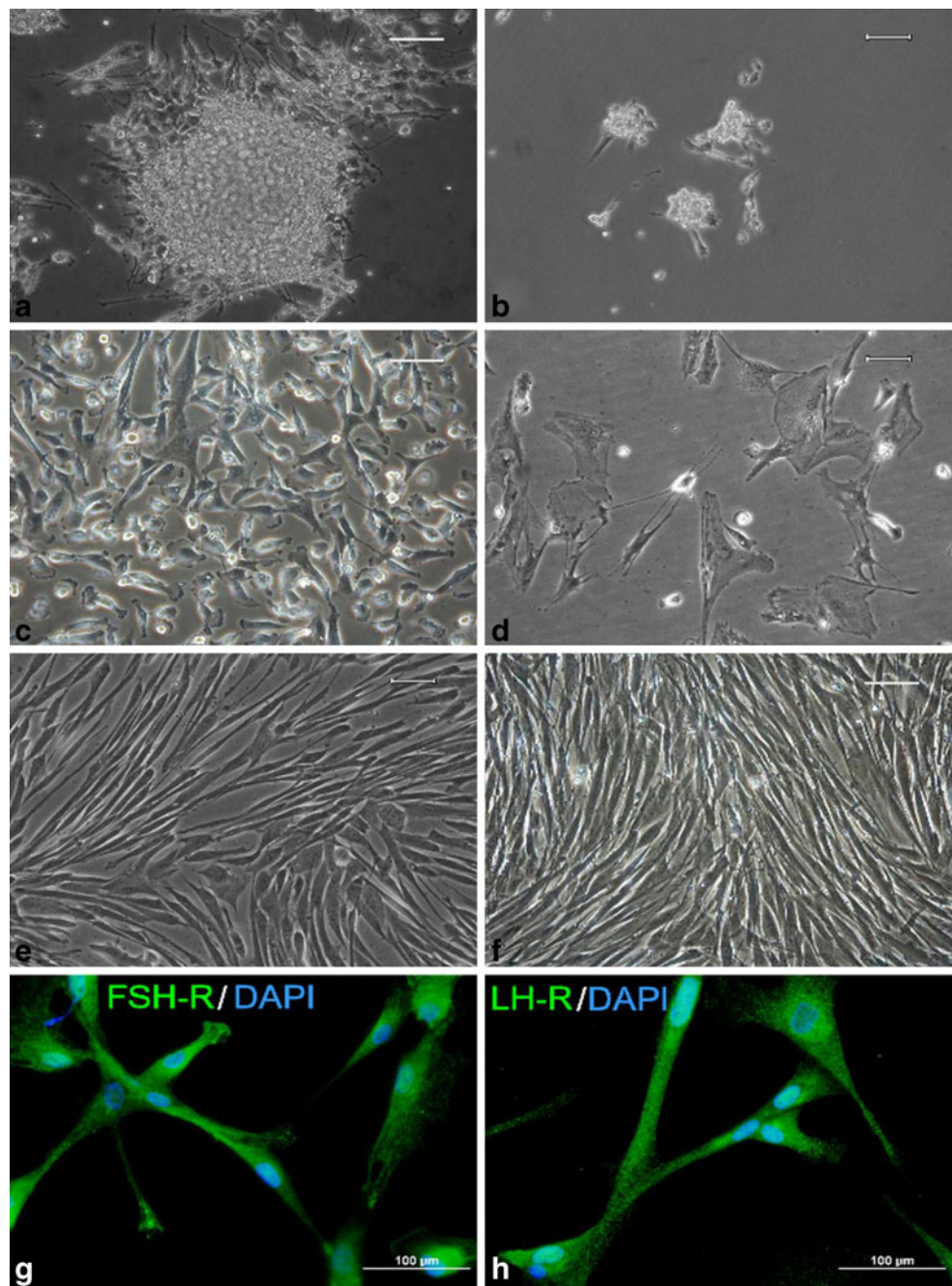
## Results

### Culture of human GCs in medium with and without FF

We evaluated the effect of follicular fluid (FF) on the prolonged survival of GCs in vitro. Isolated luteinizing GCs from each patient were splitted into two groups and cultured separately. One group was initially cultured in a medium supplemented with FF (Fig. 1a, c), whereas the other group was cultivated without FF (Fig. 1b, d). The luteinizing GCs cultured without FF consistently died within 2 weeks (Fig. 1d), whereas those initially cultivated in medium supplemented with FF remained viable for up to 1.5 months and could be passaged (Fig. 1e). In one sample, cells remained viable for up to 3.5 months (passage no. 25), Fig. 1f. These unique long-term proliferating cells were labeled as the GC8 line and became the subject of our



**Fig. 1** Representative figures of human granulosa cells (GCs) grown in 2% FCS containing medium supplemented with EGF, bFGF and FSH. **a** GCs was initiated by the addition of follicular fluid into the cultivation medium (day 1). The follicular fluid created a protein surface on the bottom of the cultivation flask that facilitated cell adhesion. **b** Morphology of GCs cultivated without follicular fluid (day 1). **c** Figure shows a small, rapidly dividing GCs with high proliferative potential, which were initially cultivated with follicular fluid (day 10). **d** Culture of GCs cultivated without follicular fluid and undergoing the cell degradation process, containing fewer dividing cells (day 10). **e** Typical spindle-shaped morphology of human GCs initially cultivated with follicular fluid (day 31). **f** Spindle-shaped cells were not morphologically changed during long-term cultivation; granulosa cells (GC8 line) initially cultivated in media with follicular fluid (day 99). Scale bar 50  $\mu\text{m}$ . **g** Immunocytochemical analysis of GCs for FSHR (day 69) and **h** LHR after initial cultivation with FF (day 69). Cells stained with the omission of a primary antibody were used as negative controls. Scale bar 100  $\mu\text{m}$



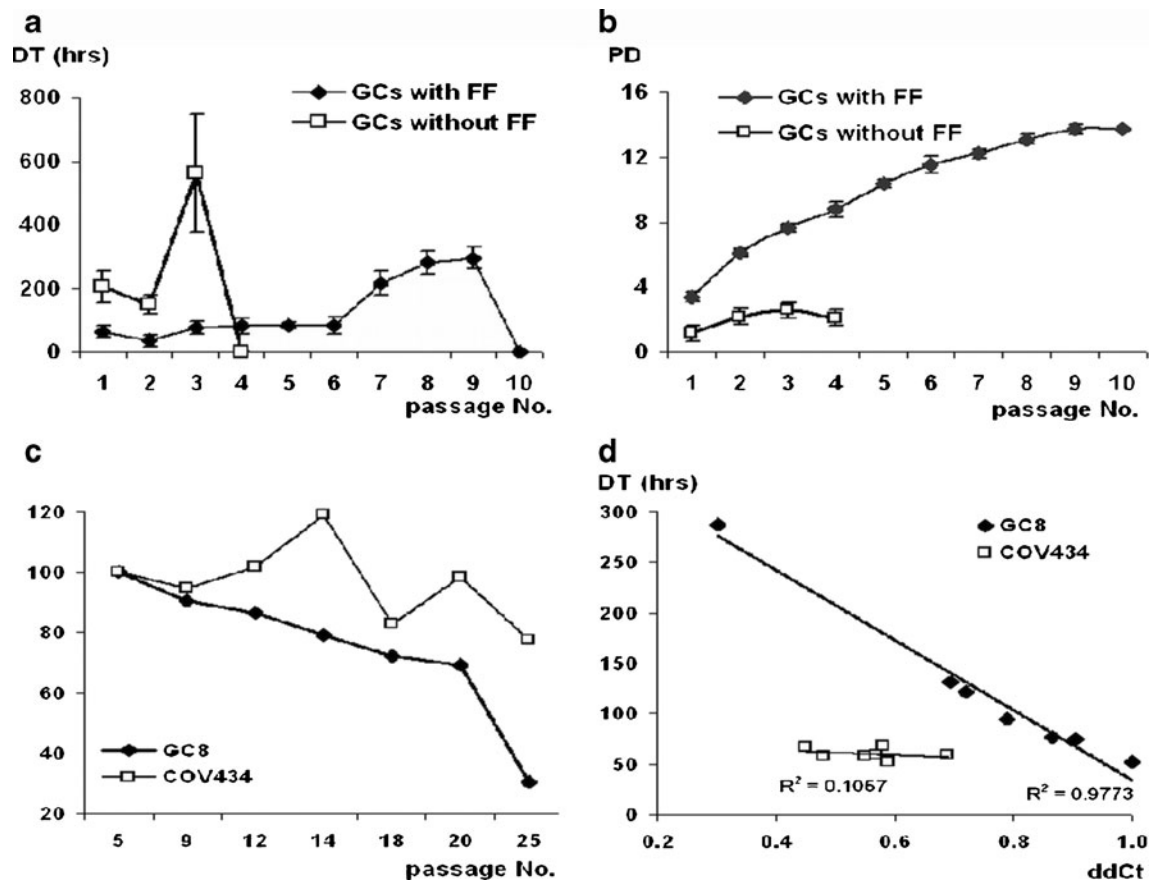
interest. Cultivated GCs were characterized by immunocytochemistry as luteinizing GCs through their expression of both FSHR (Fig. 1g) and LHR (Fig. 1h).

FF was added for the initial day of *in vitro* cultivation only. Following the addition of FF, GCs retained their morphology, established intercellular connections, and became strongly attached to the culture dish. Most of the erythrocytes were very gently washed from the protein layer of FF with PBS and large vaginal epithelial cells were not observed after trypsinization in the subculture.

The average cell viability of proliferating cells was  $90 \pm 5\%$ . After 15 days (passage no. 3), the GCs became

spindle-shaped (fibroblast-like cells). The proliferative activity of these cells was different for each patient. After 25–45 days, cells entered to either senescence or irreversible apoptotic changes leading to cell death. In contrast, GCs cultivated in the absence of FF became less viable ( $46 \pm 8\%$ ) after 7 days and lost intercellular connections.

An effect of FF on doubling time (DT) and population doublings (PD) of GCs *in vitro* is shown in Fig. 2a, b. The average DT of GCs initially cultured with FF was  $71 \pm 15$  h, whereas DT of GCs cultivated without FF was  $305 \pm 81$  h. GCs cultured without FF died after 3 passages, therefore, cumulative PD decreased, whereas those cultivated in



**Fig. 2** **a, b** GCs cultivated in medium with FF were viable for up to the ninth passage, demonstrated lower DT and a higher number of PDs. In contrast, GCs cultured in the absence of FF rapidly increased DT, slightly increased the number of PDs (cumulative data), cell died after the third passage. **c** Telomere length measurement (y-axis;%) in

long-term growing cells (GC8 line) and GC-derived tumor cell line COV434. The GC8 line showed a significant decrease in the relative telomere length (y-axis;%) with each increasing passage number ( $P < 0.01$ ). **d** Changes in doubling time (DT) versus relative telomere length (ddCt) were significant ( $P < 0.0001$ )

medium with FF remained proliferating for up to the ninth passage.

Doubling time (DT) and population doublings (PD) of GCs, GC8 and tumor granulosa cells COV434

GCs initially stimulated by FF were used for DT and PD analyses. The average DT of GCs was  $159 \pm 24$  h (mean  $\pm$  SD,  $n=24$ ). All GCs demonstrated similar proliferation potential until the third passage. Thereafter, progressive prolongation of DT was seen in the majority of GCs (24 out of 25). After the ninth passage, DT of GCs increased to 260 h and cells died. In contrast, the long-term growing GCs (GC8) were maintained until the twenty-fifth passage as presented in Fig. 3a. The average DT of the GC8 line was  $105 \pm 21$  h (mean  $\pm$  SD,  $n=3$ ) which is significantly shorter than the DT of other GCs but longer than the DT of tumor cells COV434 where the DT was  $59 \pm 3$  h.

Similarly, the cumulative PD was highest ( $44 \pm 0.3$ ) in the immortalized tumor line COV434. GCs lines increased the

PD until the ninth passage, cumulative PD was  $10 \pm 0.7$ . Contrary to this, the unique GC8 line grew to the twentieth passage, cells reached  $35 \pm 0.3$  population doublings (Fig. 3b).

In both graphs (Fig. 3a and b), we can compare the growth patterns of long-term proliferating cells (GC8) resembling the tumor cells (COV434) rather than GCs.

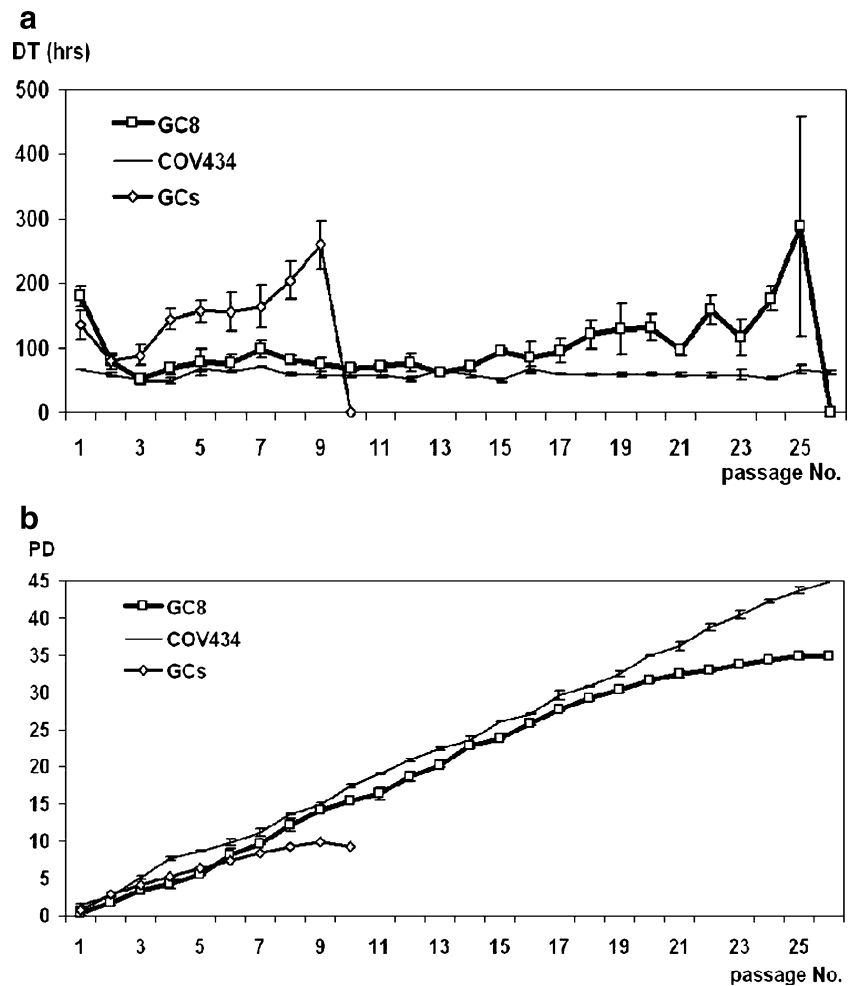
#### Karyotyping

All cultured GCs had normal karyotype in early passages. Long-term growing cells (GC8) were found to be cytogenetically stable even after 25 passages.

#### Telomere length measurement in GC8 and COV434

GC8 and COV434 included in this analysis were harvested from passages no. 5 to no. 25, i.e. they reached 4 to 40 population doublings. The GC8 line showed an overall and significant decrease in relative telomere length with each

**Fig. 3 a** The comparison of the average doubling time (DT) of GCs ( $n=24$ ), the GC-derived tumor cell line COV434 (experiment carried in triplicates) and the long-term growing granulosa cell line GC8 (experiment carried in triplicates) that were isolated and cultivated in the same way as other GC lines but presented a different characteristics. **b** Comparison of the average cumulative population doublings (PD) of GCs ( $n=24$ ), GC8 ( $n=3$ ) and the GC-derived permanent tumor cell line COV434—cumulative data



increasing passage number ( $P<0.01$ ). Also, changes in DT versus relative telomere length (ddCt) were significant ( $P<0.0001$ ) which demonstrated the prolongation of DT in cells with shortened telomere length (Fig. 2c, d).

**Flow cytometry analysis**

GCs primed with FF were used for flow cytometric analyses. In the absence of FF, the cell loss was very high and there were not enough GCs left for analysis. A phenotype of GCs lines was analyzed ( $n=25$ ) in passage no. 5. A phenotype of long-term growing cell line—GC8 was assessed in passages no. 5, 9, 12, 15 and 17. For a comparison, granulosa tumor cell line COV434 was also analyzed in passages no. 5, 9, 12, 15 and 17.

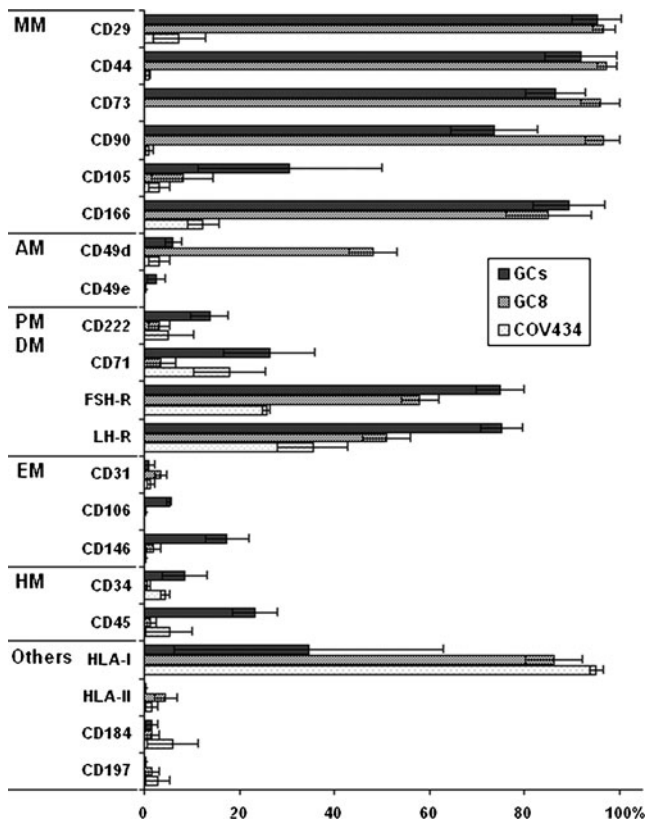
The GCs resulting data from flow cytometric analysis were as follows (Fig. 4): high expression of mesenchymal markers CD29, CD44, CD73, CD90, CD166 and gonadotropin receptors LHR and FSHR. Surface cell markers CD31, CD34, CD49d, CD49e, CD106, CD184, CD197, HLA-II were not expressed and a low expression was observed in CD45, CD71, CD105, CD146, CD222, HLA-I.

Flow cytometric analysis revealed that long-term proliferating cells GC8 (in passages no. 5, 9, 12, 15 and 17) did not express either haemopoietic markers CD34, CD45, CD184 or endothelial cell markers CD31, CD106, CD146; a low expression of CD105 was observed. Contrary to this, the cell line GC8 expressed high levels of mesenchymal markers (CD29, CD44, CD73, CD90, CD166) and HLA-I. Adhesion molecules CD49e, CD222; proliferation marker CD71; chemokine CD197 and HLA-II were not expressed and a moderate expression was observed in alfa4 integrin CD49d. A moderate expression of specific markers for GCs FSHR and LHR was observed. The exception was found in passage no. 5, where the GC8 line expressed CD71 (++) , CD105 (+) and CD222 (+) and within the following passages (no. 9, 12, 15 and 17) the expression of these surface markers was lost.

The resulting data of COV434 were as follows: high expression of HLA-I, low expression of FSHR, LHR, CD71 and CD166 and no expression of CD29, CD31, CD34, CD44, CD45, CD49d, CD49e, CD73, CD90, CD105, CD106, CD146, CD184, CD197, CD222, HLA-II.

Surprisingly, the GC8 cell line expressed high levels of HLA-I, similarly to COV34 but in contrast to other GC





**Fig. 4** Phenotypic analysis of GCs, the long-term growing granulosa cell line GC8 and the GC-derived tumor cell line COV434. Classification: <10% no expression; 10–40% low expression; 40–70% moderate expression; >70% high expression. Percentage of the population GCs (mean±SD,  $n=24$ ); GC8 (mean±SD,  $n=3$ ), COV434 (mean±SD,  $n=3$ ). Mesenchymal markers (MM), adhesion molecules (AM), proliferation markers (PM), differentiation markers (DM), endothelial markers (EM) and haemopoietic markers (HM)

lines. The unique feature of the GC8 cell line was an elevated expression of the adhesion molecule CD49d.

## Discussion

In this work, a long-term cultivation protocol of the luteinizing granulosa cell line has been established and tested. The protocol utilises our previous pilot study (16) where the addition of FF to GC improves their attachment and proliferation of primary culture.

The development of GCs is uniquely controlled by follicle stimulating hormone (FSH) during folliculogenesis, causing them to proliferate and subsequently differentiate and contribute to the formation of the follicular antrum, secreting fluids, ions, and proteins characteristic of the FF (17). FSH and LH are the primary survival factors for ovarian follicles, the antiapoptotic effects of these gonadotropins are probably mediated by the production of ovarian growth factors (18). Some of growth factors such as

fibroblast (bFGF) and epidermal growth factor (EGF) prevent apoptosis in antral follicles (19) and have potent mitogenic effects on human GCs (20). Therefore GCs were cultivated in DMEM/F12 medium supplemented with 2% fetal calf serum (FCS). In vitro proliferation of GCs was supported by FF as well as FSH and growth factors (bFGF and EGF).

Some authors used media containing 10% (or higher) concentrations of FCS for the cultivation of GCs (2, 21–23). Cultures of GCs grew to confluence through four passages but grew slowly, taking approximately 2 weeks to double in number (2). GCs plated on tissue culture plastic become apoptotic and/or undergo rapid cell death, whereas GCs plated on ECM (extracellular matrix) survive for several days in culture (24, 25). We were able to cultivate for up to the ninth passage (45 days), taking approximately 6 days to double in number. Serum contains multiple factors (polypeptides, hormones, growth factors and cytokines, or binding proteins soluble receptors) that might influence cell function (1) and GC luteinization is accelerated in medium supplemented with serum (26). FF is as poorly defined as any FCS used in cultivation media. However samples of FF were obtained from ovarian follicles together with GCs always from same patient. FF is the intercellular environment crucial for fertilization and early embryonic development, as the follicle enlarges a fluid-filled antrum forms due to the replication of the GCs (3).

The crucial difference between our approach and other approaches was the application of follicular fluid. FF facilitates the communication of the oocyte with the surrounding environment leading to oocyte maturation accompanied by a differentiation and proliferation of GCs in vivo. It is an indicator of secretory activities and metabolism of GCs (27). Pioneering work suggested that GCs of mammalian follicles synthesize glycosaminoglycans at the time of antrum formation and secrete them into the FF (28). These glycosaminoglycans were identified as chondroitin sulphate, dermatan sulphate, hyaluronan and a heparin. GCs cultured in vitro synthesize and secrete proteoglycans with chemical and physical properties very similar to those isolated from FF. Proteoglycans are macromolecules formed by a protein backbone to which one or more glycosaminoglycan side chains are co-valently attached. They are involved in matrix formation, cell-cell and cell-matrix adhesion, cell proliferation and migration, and show co-receptor activity for growth factors (29). Our results suggest that the nutrients in a medium with addition of FF are essential to maintain the proliferative potential of primary culture of GCs.

Several studies have shown that the in vitro culture of GCs plated on matrices of ECM proteins of various types and densities leads to changes in cell adhesion



and shape (25). Normal GC morphology *in vivo* is round, yet GCs seeded onto uncoated tissue culture plastic resemble fibroblast-like shape, flattened in a monolayer and spread with little interaction between neighboring cells (24, 25, 30, 31). When plated on type I collagen gels, isolated basal lamina, or Matrigel (BD, USA), however, GCs retain their spherical, epithelioid shape (25, 30, 31). Our results suggest that adhesion of individual cells and also clumps of GCs were initiated by the addition of FF into the cultivation medium (Fig. 1a,c). FF partially precipitated and created a thin layer on the bottom of the cultivation plate which facilitated the cell adhesion. GCs retained their round morphology, constructed intercellular connections, and became strongly attached to the culture dish. We observed that cells were in clumps of small rounded cells (Fig. 1a), after trypsinization they took an epithelial-like appearance (Fig. 1c). After approximately 7–15 days (Fig. 1e), GCs became spindle-shaped (fibroblast-like cells) because GCs were seeded onto uncoated tissue culture plastic without FF. Similar changes in morphology cultivated GCs were observed in other works (2, 32). Kossowa-Tomaszczuk cultivated GCs in monolayers, cells invariably became luteinized and converted from epithelial-like into a fibroblast-like morphology, explaining why the latter morphology became dominant during prolonged culture (32). Gutiérrez et al. published that the spindle-shaped GCs may have lost aromatase activity, cytoplasmic changes compatible with luteinization (33).

The addition of FF in culture medium showed an overall and significant increase in GC proliferation (Fig. 2a, b), viability and cell survival. Our work brings the first evidence of the determination of population doublings (PD) and doubling time (DT) of prolonged cultivated GCs initiated by the addition of FF. The progressive prolongation of DT was seen in the majority of GCs until the ninth passage (days 25–45). In contrast, unique long-term growing cells (GC8) isolated from the patient remained viable for up to passage no. 25 (day 102, Fig. 1f).

The average DT of all GCs was  $159 \pm 24$  h versus the DT of the long-term growing GC8 where DT was lower— $105 \pm 21$  h. All GCs demonstrated a similar PD trend until the ninth passage. All GCs lines stopped growing in the ninth passage because cells died. Only the long-term growing line GC8 increased its PD in the twenty-fifth passage (Fig. 3a, b). The total cumulative PD of all GCs was  $10 \pm 0.7$  while the GC8 line reached  $35 \pm 0.3$  PD. Immunofluorescent microscopy showed that all cultured granulosa cells (GC8 also) retain their FSH receptor expression through all passages (Fig. 1g, h).

The population of GCs in a healthy follicle is not uniform but rather consists of subpopulations of differ-

entiated and less differentiated cells, the latter being more capable of mitosis (10). We suggest that we isolated a population of granulosa cells (GC8), which contained a subpopulation of less differentiated cells. Immunofluorescence microscopy showed that these cells also express stem cell marker nestin (results not shown). For the first time in 2009, Kossowa-Tomaszczuk et al. demonstrated that prolonged culture of luteinizing GCs in medium supplemented with LIF (leukemia-inhibiting factor) allows the selection of less differentiated GCs, which exhibited a certain degree of plasticity and multipotency. FF contains a high concentration of LIF which stimulates GCs through LIF-R physiologically in mature follicles (32). FF was added for the initial day of *in vitro* cultivation however its influence was evident during the whole cultivation. There is the possibility of an alternative mechanism, i.e. de-differentiation of granulosa cells (GC8) *in vitro*. Currently we have no evidence for this mechanism.

A further explanation for the long growing GC8 could be the neoplastic transformation within *in vitro* cultivation. This was the reason for using the standard tumor cell line COV434 to compare with GCs. Zhang et al. published that during 38 passages the properties of the COV434 tumor granulosa cells did not undergo significant changes (13). Throughout the *in vitro* cultivation of tumor cells COV434, stable DT in every passage was observed. The gradually increasing PD was the highest in tumor line COV434 where growth continued even after 25 passages. Growth patterns of the GC8 line were closer to tumor cells COV434 than to other GCs (Fig. 3a, b). Therefore, karyotype was analyzed to confirm GCs chromosomal stability and to exclude neoplasm transformation. All cultivated GCs as well as the long-term growing cells GC8 showed normal karyotype and chromosomal stability in early passages (GC8 also after reaching twenty-fifth passage).

GCs proliferation potential was furthermore analyzed by the measurement of the length of the telomere repeat sequence. Expression of telomerase, the enzyme that maintains telomere length, is repressed in most adult differentiated somatic cells, although it is active in human ovaries and testes (34), in the endometrium (35), in stem cells present in the bone marrow and blood cells (36), in the liver (37), in the epidermis (38), and all structures characterized by an intense proliferative status. In fact, the introduction of telomerase into normal human somatic cells has been shown to extend normal cell life by ~20 doublings (39). Thus, telomerase activity is present in regenerative tissues with high proliferative need, where telomere loss would deplete the stem cell populations (3). We compared the length of telomeres of long-term growing cells (GC8) and immortal tumor granulosa cells COV434, the maintenance of the telomere length seems to guarantee cell immortality. We found that changes in

doubling time (DT) versus relative telomere length (ddCt) of the GC8 line were significant (Fig. 2d) which demonstrated the prolongation of DT in cells with a shortened telomere length.

Moreover, GCs were characterized by the determination of the phenotype and its comparison with the GCs-derived tumor cell line COV434 (Fig. 4). Based on current knowledge of human stem cell phenotype, twenty-one surface markers were chosen and determined by flow cytometry. GCs originate from mesoderm, therefore especially mesenchymal markers were examined. Analysis revealed that GCs (in passage no. 5,  $n=24$ ) expressed high levels of the mesenchymal markers CD29, CD44, CD73, CD90, CD166 and low levels of the marker CD105. Kossowa-Tomaszcuk et al. published similar results—CD29, CD44, CD90, CD105, CD166, but not CD73, were expressed by substantial subpopulations of the freshly collected GCs (32). Several recent studies, using techniques of molecular biology, confirmed that the GCs are the only cell type expressing the FSH-R, thus expression of the FSH-R is strictly gonad- and highly cell specific (40). LH-R expression can be demonstrated in a variety of organs and tissues (41). These markers, FSH-R and LH-R were highly expressed on GCs in vitro. The typical marker for haematopoietic cells, CD45, was present in low concentration probably due to contamination with blood cells in early passages, CD34 and CD184 were not expressed.

Moreover, due to a different phenotypic pattern, we suggest the presence of several subpopulations of GCs as in preovulatory follicles, each expressing different markers according to their degree of differentiation. For example, the GC8 line remained viable for up to the twenty-fifth passage and presented an increased expression of adhesion marker integrin CD49d and HLA-1, cells lost expression of markers CD45, CD71, CD105, CD146, CD222 and also decreased the expression of markers (FSH-R, LH-R) during in vitro cultivation. It is noteworthy that the granulosa cell (GC8) showed elevated expression of the alpha4 integrin subunit (CD49d). Integrins comprise a family of cell surface receptors that mediate cell-ECM and cell-cell adhesions by interacting with ECM proteins such as fibronectin, laminin, collagen and vitronectin, as well as with counterreceptors of the immunoglobulin superfamily of cell adhesion molecules (42). At a cellular level, integrins are involved in a number of cellular processes such as cell proliferation and differentiation, cell migration, cytoskeletal organization and cell polarization. (43).

We hypothesize that long-term proliferating cells (GC8) expressing different markers were less differentiated subpopulation isolated from ovarian follicle. Moreover, the

expression of the proliferation marker CD71 (Transferrin-R) was higher in the GC8's fifth passage together with the expression of the mesenchymal marker CD105 and CD222 (IGF-II R). It is possible that these three markers could be expressed differently in several subpopulations of GCs in preovulatory follicles.

Isolated GCs lines with well defined biological characteristics may be used for the development of in vitro maturation of immature oocytes. We characterized the long-term cultivated GCs obtained from patients and we compared them with GCs-derived permanent tumor cell line COV434. The results of the proliferation and phenotype analyses demonstrated that there is a fundamental difference between the GCs from patients and the GCs-derived cell line COV434. Significantly different phenotypes of GCs-derived tumor cell line COV434 was determined and we found no expression of mesenchymal markers CD29, CD44, CD73, CD90, CD105, CD166 and low expression of FSHR, LHR, CD166 (Fig. 4). Thus the in vitro model of GCs-derived permanent tumor cell line COV434 can be inconvenient because cells were neoplasm transformed. On the other hand, GCs obtained during the collection of oocytes from patients could represent a suitable In vitro model.

Our research has shown that some cells of ovarian follicle origin possess stem cell-like characteristics of the mesenchymal lineage. Since these cells present the FSH-R on their surface, it is likely that they are derived from the GCs. These ovarian stem cells can be cultured over prolonged time period while maintaining the biological characteristics of GCs. The long-term culture of those GCs can be used in studies of oocyte in vitro maturation, in differentiation studies of embryonic stem cells co-cultured with GCs or simply to study the endocrine functions of GCs in vitro.

## Conclusions

Our cultivation protocol enables GCs to proliferate for 45 days and up to 100 days in the case of the GC8 cell line. This new approach was due to the addition of follicular fluid to the cultivation medium with a low serum concentration. Doubling time and population doublings assessment demonstrated the possibility of “long-term” cultivation of isolated GCs. Primary culture of human GCs initiated by the addition of FF could represent a suitable in vitro model to study disruptions of hormonal stimulation and serious complications of assisted reproduction protocols.

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