



# All that glitters is not gold: a stereological study of human donor oocytes

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## Research Article

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### Summary

Here we report a quantitative analysis of human metaphase II (MII) oocytes from a 22-year-old oocyte donor, retrieved after ovarian-controlled hyperstimulation. Five surplus donor oocytes were processed for transmission electron microscopy (TEM), and a stereological analysis was used to quantify the distribution of organelles, using the point-counting technique with an adequate stereological grid. Comparisons between means of the relative volumes ( $V_v$ ) occupied by organelles in the three oocyte regions, cortex (C), subcortex (SC) and inner cytoplasm (IC), followed the Kruskal–Wallis test and Mann–Whitney  $U$ -test with Bonferroni correction. Life cell imaging and TEM analysis confirmed donor oocyte nuclear maturity. Results showed that the most abundant organelles were smooth endoplasmic reticulum (SER) elements (26.8%) and mitochondria (5.49%). Significant differences between oocyte regions were found for lysosomes ( $P = 0.003$ ), cortical vesicles ( $P = 0.002$ ) and large SER vesicles ( $P = 0.009$ ). These results were quantitatively compared with previous results using prophase I (GV) and metaphase I (MI) immature oocytes. In donor MII oocytes there was a normal presence of cortical vesicles, SER tubules, SER small, medium and large vesicles, lysosomes and mitochondria. However, donor MII oocytes displayed signs of cytoplasmic immaturity, namely the presence of dictyosomes, present in GV oocytes and rare in MI oocytes, of SER very large vesicles, characteristic of GV oocytes, and the rarity of SER tubular aggregates. Results therefore indicate that the criterion of nuclear maturity used for donor oocyte selection does not always correspond to cytoplasmic maturity, which can partially explain implantation failures with the use of donor oocytes.

### Introduction

According to data published by the World Health Organization, infertility is a condition that affects ~48 million couples and 186 million individuals worldwide (WHO, 2020). Female infertility is responsible for approximately half of the cases, with ovarian ageing being a current societal dilemma (Walker and Tobler, 2022).

Oocyte quality is a critical component in overall fertility, as oocyte competence is critical for normal fertilization, embryonic development, implantation and full pregnancy (Conti and Franciosi, 2018; Sirait *et al.*, 2021). Oocyte competence is acquired during oogenesis, where the oocyte passes through a sequence of maturation changes involving both nuclear and cytoplasmic domains (Arroyo *et al.*, 2020). Nuclear maturation includes a cascade of events culminated by the pre-ovulatory peak of luteinizing hormone (LH), which allows the oocyte to resume meiosis and ovulate (Sánchez and Smitz, 2012). This process is characterized by nuclear envelope breakdown (GVBD: germinal vesicle breakdown) in prophase I oocytes (GV oocytes), followed by the condensation and alignment of chromosomes in a metaphase plate-I (MI oocytes). MI oocytes then enter the first meiotic division, with extrusion of the first polar body (PB1), containing half of the chromosomes, and condensation and alignment of the other half of the chromosomes in a metaphase plate-II (MII oocytes), with oocytes remaining in a MII arrest until fertilization (Arroyo *et al.*, 2020). Nuclear maturation must occur in coordination with cytoplasmic maturation (Eppig, 1996). Cytoplasmic maturation is characterized by the

accumulation of mRNA, proteins, substrates and nutrients (Watson, 2007; Richani and Gilchrist, 2018; Cornet-Bartolomé *et al.*, 2021), as well as by the remodelling of organelles and the cytoskeleton (Mao *et al.*, 2014; Coticchio *et al.*, 2015; Reader *et al.*, 2017), in an intimate cross-talk with the surrounding follicular cells (Dumesic *et al.*, 2015; May-Panloup *et al.*, 2016).

Although the objective in assisted reproductive treatments (ART), using controlled hormonal ovarian hyperstimulation, is to obtain a maximum number of MII oocytes, in ~10–15% of the cycles immature oocytes, GV and MI, are also retrieved (Ebner *et al.*, 2003; Pinto *et al.*, 2009; Braga *et al.*, 2020), and, in a lower percentage, oocytes may display cytoplasmic and extracytoplasmic dimorphisms (Rienzi *et al.*, 2011; Sá *et al.*, 2011; Braga *et al.*, 2013, 2020; Sousa *et al.*, 2015, 2016a; Stigliani *et al.*, 2018), due to maturation defects (Windt *et al.*, 2001; Tripathi *et al.*, 2010; Conti and Franciosi, 2018; Ozturk, 2020). Some immature oocytes may resume meiosis during *in vitro* culture, but show reduced developmental competence (Ebner *et al.*, 2003, 2006; Shu *et al.*, 2007; Azevedo *et al.*, 2014; De Vos *et al.*, 2021; Karavani *et al.*, 2021).

Oocyte competence can be evaluated through morphologic and molecular analyses. The ultrastructural morphology of human oocytes at different maturation stages was evaluated in oocytes obtained from ART cycles (Baca and Zamboni, 1967; Zamboni *et al.*, 1972; Sathananthan and Trounson, 1982, 2000; Sathananthan, 1985, 1994, 1997, 2003, 2013; Sundström *et al.*, 1985; Motta *et al.*, 1988, 2000; Familiari *et al.*, 2006; Sathananthan *et al.*, 2006; Morimoto, 2009; Nottola *et al.*, 2014). Qualitative morphological studies have also been recently conducted in donor oocytes (Trebichalská *et al.*, 2021). Molecular research involved the analysis of oocyte and cumulus cells transcriptome (Cillo *et al.*, 2007; Gasca *et al.*, 2007; Wells and Patrizio, 2008; Ouandaogo *et al.*, 2012; Coticchio *et al.*, 2017), proteome (Virant-Klun *et al.*, 2016; Shen *et al.*, 2017) and metabolome (Venturas *et al.*, 2021), or zona pellucida gene mRNA expression (Canosa *et al.*, 2017), with development of oocyte quality markers (Sirait *et al.*, 2021).

Despite the importance of descriptive morphological studies, the qualitative evaluation of oocyte morphological characteristics can be incomplete or erroneous. As knowledge of the real morphology of the human oocyte is critical for the clinical development of new methods of cryopreservation and *in vitro* maturation, Pires-Luis *et al.* (2016) and Coelho *et al.* (2020) conducted pioneer quantitative studies with a stereological approach in GV and MI oocytes, respectively. In these studies, the relative volumes of the different organelles were measured and the distribution of organelles in the different regions of the oocyte, cortex, subcortex and inner cytoplasm (IC), were evaluated.

At this time, the use of donor oocytes showed an exponential increase due to the large improvements in the cryopreservation technique and luteal phase endometrium preparation (Walker and Tobler, 2022). Although with the use of donor oocytes live birth rates are high, there is still a place for implantation failures. This has been attributable to recipient uterine conditions or partner male factors [European IVF-Monitoring Consortium (EIM) for the European Society of Human Reproduction and Embryology (ESHRE), 2021; Sciorio *et al.*, 2021; Williams *et al.*, 2022], with the competence of donor oocytes being admitted to be optimized.

To our knowledge, this ultrastructural study is the first for donor MII oocytes using a stereological approach, in which we intended to quantify the spatial distribution of organelles.

Surprisingly, we found a set of donor oocytes showing nuclear maturity but cytoplasmic immaturity. The results of the present report therefore provide evidence that even donor oocytes can show impaired competence.

## Materials and methods

### Ethical approval

Ethical guidelines were followed when conducting the research. According to the National Law on Medically Assisted Procreation (Law 32/2006) and the National Council for Medically Assisted Procreation guidelines (CNPMA, 2018), the use of clinical databases and surplus gametes for research may be used under strict individual anonymity and after patient written informed consent. Surplus donor oocytes were used after patients signed an informed consent agreeing to share their own gamete samples for the present study. The present laboratory experiments were executed under the Joint Ethics Committee of the Hospital and University, CHUP/ICBAS approval number 2019/CE/P017 (266/CETI/ ICBAS). This work did not involve human or animal experiments and therefore the provisions of the Declaration of Helsinki as revised in Tokyo 2004 do not apply.

### Patients

This research was performed on five MII oocytes recovered from a 22-year-old oocyte donor after controlled ovarian stimulation during a donation programme performed at the Centro de Procriação Medicamentada Assistida (CPMA), Centro Materno-Infantil do Norte (CMIN) Albino Aroso, Centro Hospitalar do Porto (CHUPorto), which is the National Gamete Bank.

### Ovarian-controlled hyperstimulation

The oocyte donor underwent controlled ovarian hyperstimulation with a gonadotropin-releasing hormone antagonist protocol (GnRH; MSD, Hertfordshire, UK). Recombinant follicle-stimulating hormone (rFSH; Teva Biotech, Germany) was used for ovarian stimulation. A GnRH agonist (0.2 mg triptorelin; Ipsen Pharma Biotech, Signes, France) was administered for triggering oocyte maturation, 36 h before follicular puncture (Huirne *et al.*, 2007; Pinto *et al.*, 2009).

### Gamete and embryo handling

Procedures were performed on a K-Systems laminar flow chamber with a thermal base at 37°C (Cooper Surgical, Malöv, Denmark). Cumulus–oocyte complexes (COC) were collected in 1-well culture dishes (Falcon, Corning, New York, NY, USA) with SynVivo Flush medium (without heparin; Origio, Malöv, Denmark) and their handling and culture were performed under paraffin oil (Ovoil-100, Vitrolife, Frölunda, Sweden). After collection, COCs were washed and then cultured in Sequential Fert medium (Origio) in an Esco incubator (MRI-6A10, EscoMedical, Singapore, Singapore) (37°C, 5% O<sub>2</sub>, 6% CO<sub>2</sub>, 89% N<sub>2</sub>) for 2 h. After denudation with recombinant hyaluronidase (ICSI Cumulase, Origio), oocytes were transferred to 4-well culture dishes (Nunc, Thermo Scientific, Denmark) with Sequential Fert medium (Origio) and processed for transmission electron microscopy (TEM). The remaining donor oocytes were vitrified under the National Gamete Bank programme according to Kitazato methodology (Kitazato Corporation, Japan).

### Transmission electron microscopy

Oocytes were fixed in Karnovsky solution (2.5% glutaraldehyde, 4% paraformaldehyde, 0.15 M sodium cacodylate buffer) (Sigma-Aldrich, St. Louis, USA; Merck, Darmstadt, Germany) at room temperature for 30 min, followed by 2 h at 4°C. After washing in 0.15 M sodium cacodylate buffer, pH 7.3 (Merck) for 2 h at 4°C, oocytes were post-fixed with 2% osmium tetroxide (Merck) in buffer containing 0.8% potassium ferricyanide (Merck) for 2 h at 4°C, washed in a buffer for 10 min, serially dehydrated in ethanol (Panreac, Barcelona, Spain), equilibrated with propylene oxide (Merck), and embedded in Epon (Sigma). Semithin and ultrathin sections were cut using diamond knives (Diatome, Hatfield, Switzerland) in an LKB ultramicrotome (Leica Microsystems, Wetzlar, Germany). A random number table was used to select the initial cut, and then oocytes were serially sectioned and sampled every 10 µm. Ultrathin sections were collected on 100 mesh formvar carbon-coated copper grids (Taab, UK) and stained with 3% aqueous uranyl acetate for 20 min (BDH, Poole, UK) and Reynolds lead citrate for 10 min (Merck) at room temperature in a light-protected environment. Ultrathin sections were observed on a JEOL 100CXII transmission electron microscope (JEOL, Tokyo, Japan) at 60 kV (Sousa and Tesarik, 1994; El-Shafie *et al.*, 2000).

### Stereological and statistical analysis

Systematic sampling was performed on each microscope grid, with photographs taken at alternate TEM field spaces when more than 50% of the field was occupied by the oocyte cytoplasm. Images were taken at ×5300 magnification and printed at 20.2 cm × 20.2 cm. A classical manual stereological technique based on point counting with a suitable stereological grid was used. The grid was placed over the printed photographs and the number of grid points placed over each organelle was noted. The relative volume ( $V_v$ ) of each organelle was obtained by applying the formula  $V_v(\text{organelle, oocyte}) = [\text{number of dots (organelle)}/\text{number of dots (oocyte)}] \times 100$  (%) (Weibel *et al.*, 1966).

Organelles included in the present evaluation were: cortical vesicles (CV), mitochondria, dictyosomes, lysosomes, medium-sized vesicles containing granular materials (VZP), smooth endoplasmic reticulum (SER) small vesicles (SER-SV), SER medium vesicles (SER-MV), SER large vesicles (SER-LV), SER very large vesicles (SER-VLV), SER isolated tubules (SER-IT), SER tubular aggregates (aSERT), and total SER without VLV (SV, MV, LV, IT, aSERT).

Each oocyte was divided into three regions, from the oolemma up to the cell centre: cortex (5 µm), subcortex (5–10 µm), and IC (>10 µm) (Figure 2A). A stereological procedure was adopted, applying the formula  $V_v(\text{organelle, cortex/subcortex/IC}) = [\text{number of points (organelle)}/\text{number of points (cortex/subcortex/IC)}] \times 100$  (%) (Pires-Luis *et al.*, 2016).

Statistical analysis was performed using Microsoft® Excel 2022 and SPSS version 27.0 software (IBM Corp, Foster City, California, CA, USA). The results are presented as mean, standard error of the mean (SEM = standard deviation/ $n^{1/2}$ ), and coefficient of variation (CVar = standard deviation/mean). Normal distribution was tested using the Kolmogorov–Smirnov test. The samples did not follow a normal distribution, and therefore non-parametric tests were used. To compare the means of  $V_v$  (organelle, oocyte),  $V_v$  (organelle, cortex),  $V_v$  (organelle, subcortex) and  $V_v$  (organelle, IC) the Kruskal–Wallis test and the Mann–Whitney *U*-test with Bonferroni correction were used. The level of statistical significance was set at  $P < 0.05$ .

### Results

When observed under the inverted microscope, live MII oocytes displayed a rounded shape and a uniform fine granular cytoplasm, being surrounded by a thick translucent zona pellucida (ZP) and separated from the ZP by a short and narrow perivitelline space, where the PB1 was located (Figure 1A). In semithin sections, the oocyte evidenced the same structures, being additionally evident microvilli, CV and large vesicles. The oocyte metaphase II plate was also discernible, being placed slightly away from the cortical region. The PB1 showed chromosomes and the characteristic organelles of the cortical ooplasm (Figure 1B,C).

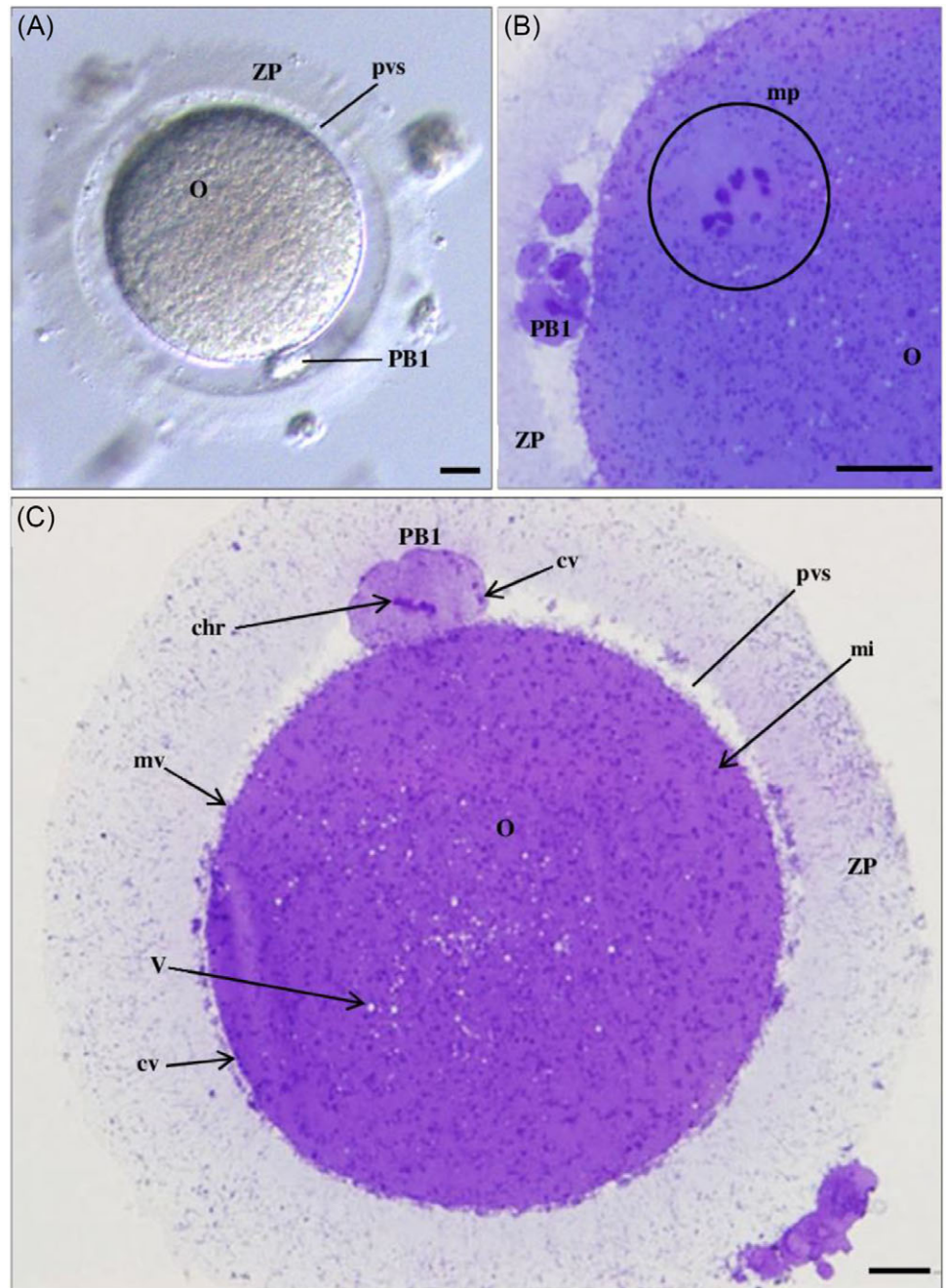
At the ultrastructural level, the oocyte surface exhibited numerous microvilli (Figure 2A,B), and in the perivitelline space the PB1 evidenced chromosomes and cortical ooplasm characteristics, including CV (Figure 2B). In the ooplasm, rows of electron-dense CV were observed under the oolemma and as isolated vesicles in the oocyte subcortex (Figure 2A). The metaphase II plate was slightly displaced towards the upper IC (Figure 2C). Mitochondria, SER-IT, SER-SV, SER-MV (Figure 2A) and SER-LV (Figure 3A) were observed in all three regions of the oocyte. The SER-VLV were found in the IC of one oocyte (Figure 3B). The aSERT was observed in all oocyte regions and also in one oocyte (Figure 4A,B). The VZP were observed in the oocyte cortex of two oocytes (Figure 4C). Lysosomes were observed in the cortex and inner ooplasm (Figure 4D). Dictyosomes were observed in the subcortex and IC of two oocytes (Figure 4E). The oocyte surface contained numerous tiny coated vesicles (Figure 4F). Multivesicular bodies, annulate lamellae, rough endoplasmic reticulum cisternae, lipid droplets and polyribosomes were not observed.

The present stereological analysis was performed on 204 photographs, with 96 representations of the cortex, 96 of the subcortex and 176 of the IC. The following organelles were quantified: mitochondria, dictyosomes, lysosomes, CV, VZP and SER elements (SER-SV, SER-MV, SER-LV, SER-VLV, SER-IT and aSERT).

The most predominant organelles in the total oocyte (Table 1) were SER-SV ( $V_v$  8.84%), SER-IT ( $V_v$  7.16%) and mitochondria ( $V_v$  5.49%). This was followed by SER-MV ( $V_v$  2.01%), LV-SER ( $V_v$  0.79%), aSERT ( $V_v$  0.61%), lysosomes ( $V_v$  0.33%), CV ( $V_v$  0.31%), SER-VLV ( $V_v$  0.30) and dictyosomes ( $V_v$  0.01%).

The most common organelles per oocyte region were (Table 1): in the cortex, SER-SV ( $V_v$  9.67%) followed by SER-IT ( $V_v$  8.19%) and mitochondria ( $V_v$  5.66%); in the subcortex, SER-SV ( $V_v$  8.74%), SER-IT ( $V_v$  7.22%) and mitochondria ( $V_v$  5.79%); and in the IC, SER-SV ( $V_v$  8.40%), SER-IT ( $V_v$  7.2%) and mitochondria ( $V_v$  5.33%).

The three oocyte regions were compared using the Kruskal–Wallis test (Table 2) and no significant differences were found for mitochondria ( $P = 0.677$ ), dictyosomes ( $P = 0.317$ ), SER-SV ( $P = 0.677$ ), SER-MV ( $P = 0.102$ ), SER-VLV ( $P = 0.368$ ), SER-IT ( $P = 0.336$ ) and aSERT ( $P = 0.980$ ). Conversely, significant differences were found for lysosomes ( $P = 0.003$ ), CV ( $P = 0.002$ ), VZP ( $P = 0.032$ ) and SER-LV ( $P = 0.009$ ). Strict pairwise comparisons for the three regions using the Mann–Whitney *U*-test, Bonferroni corrected (Table 2), showed that there were significant differences in the  $V_v$  of CV between the three regions of oocytes ( $P = 0.008$ ), lysosomes showed significant differences between cortex vs IC and subcortex vs IC ( $P = 0.008$ ), and SER-LV between cortex vs IC ( $P = 0.032$ ) and subcortex vs IC ( $P = 0.008$ ).



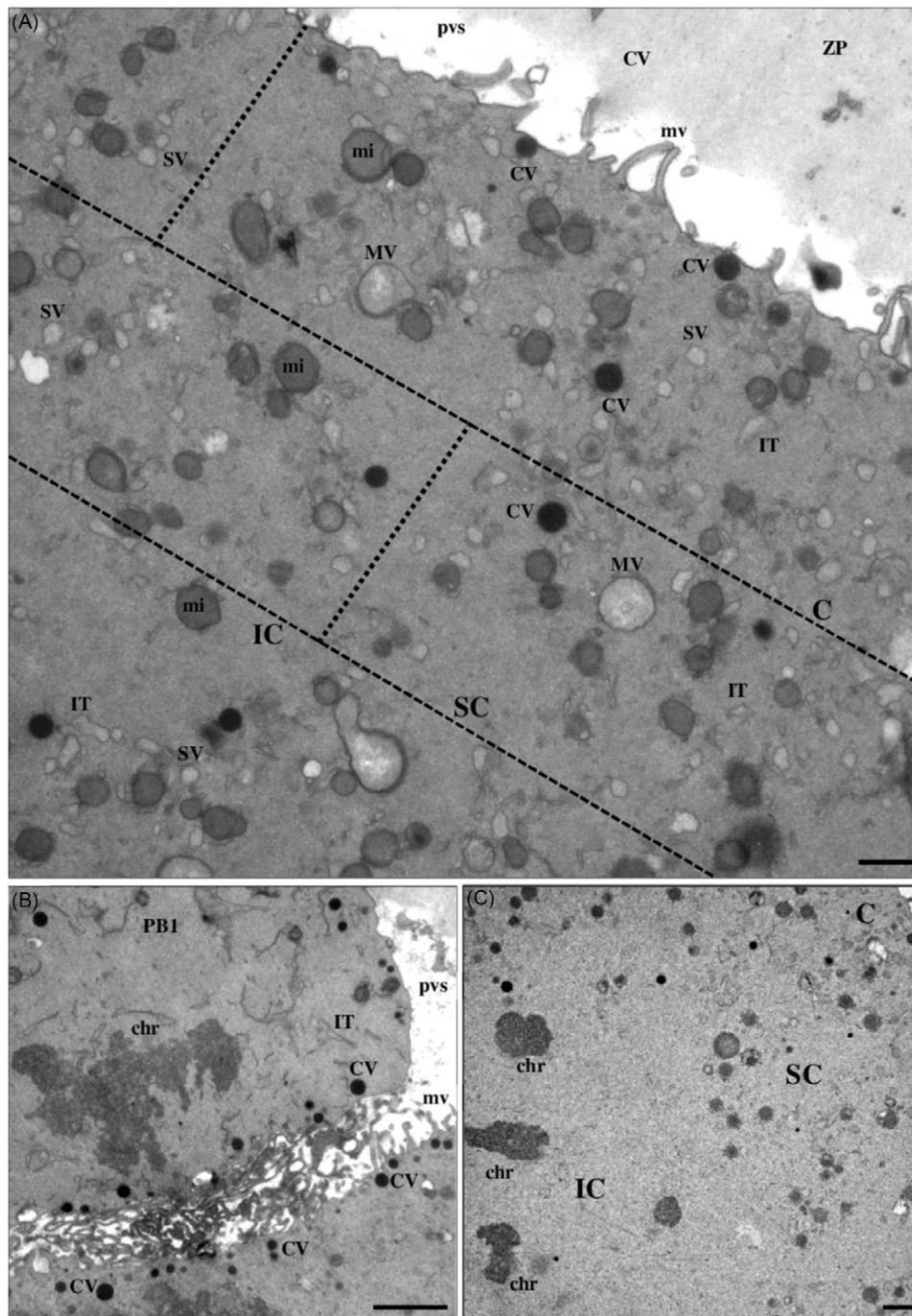
**Figure 1.** Metaphase II (MII) oocyte. (A) Live cell observed in the inverted microscope and (B, C) its correspondent semithin sections. The oocyte (O) was separated from the zona pellucida (ZP) by the perivitelline space (pvs), where resided the first polar body (PB1). In all cases, the ooplasm presented a homogeneous appearance. In semithin sections, all oocytes presented a metaphase place (mp) slightly dislocated towards the upper inner region. Chromosomes (chr) and cortical vesicles (cv) appeared evident in the PB1. The oocyte surface displayed numerous microvilli (mv). In the ooplasm, cortical vesicles (cv), mitochondria (mi) and large vesicles (V) can be observed. Bars: (A) 20  $\mu$ m; (B, C) 10  $\mu$ m.

## Discussion

Donor oocytes are an important procedure in ART and in the last 30 years have become a more prevalent therapy of choice for many infertile patients (Yeh *et al.*, 2014). This study is based on the use of donor oocytes from a young woman by a recipient woman in ART procedures. Indications for donor oocytes include advanced reproductive age, diminished ovarian reserve, recurrent poor oocyte or embryo quality in IVF attempts, hypothalamic hypogonadism, and in some cases with the possibility of transmitting a genetic defect (Taylor, 2019; Sciorio *et al.*, 2021; Williams *et al.*, 2022). In oocyte donor cycles, in which ovarian hyperstimulation is used, it is necessary that nuclear and cytoplasmic maturation occurs in a

coordinated manner to ensure optimal conditions for subsequent fertilization (Conti and Franciosi, 2018), but which does not always occur, resulting in nuclear, cytoplasmic or extracytoplasmic morphological abnormalities (Ten *et al.*, 2007; Setti *et al.*, 2021).

In previous works we presented the stereological analysis of human immature oocytes at the GV stage (Pires-Luís *et al.*, 2016) and MI stage (Coelho *et al.*, 2020). The present study was designed to apply the same technique to mature MII oocytes. To this end, we obtained 13 surplus donor MII oocytes, retrieved from six donors during oocyte donor programmes. In one case, the oocyte donor donated five oocytes for research, three cases gave two oocytes each, and two cases gave one oocyte each. In all cases, nuclear maturity was verified through inspection under an

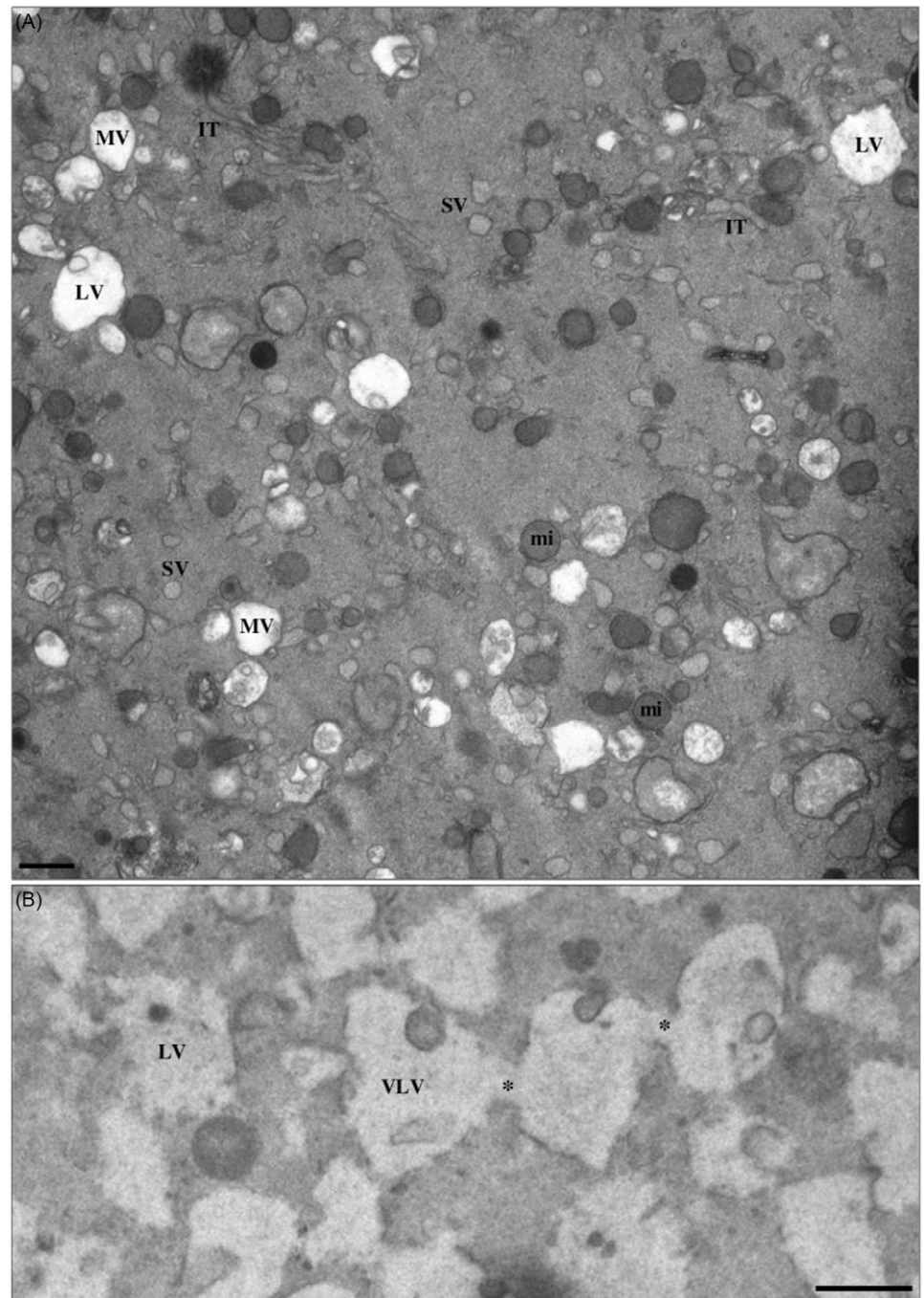


**Figure 2.** Ultrastructural images of metaphase II (MII) oocytes. (A, B) MII oocytes, showing the three regions, cortex (C), subcortex (SC) and inner (IC) cytoplasm. The oocyte surface is separated from the zona pellucida (ZP) by the perivitelline space (pvs), which contains oocyte microvilli (mv). Rows of cortical vesicles (cv) are observed under the oolemma and as isolated cortical vesicles in the cortex and subcortex regions. In the ooplasm mitochondria (mi) and smooth endoplasmic reticulum (SER) small (SV) and medium (MV) sized vesicles and isolated tubules (IT) can be observed. SER vesicles were barely associated with mitochondria. (B) The first polar body (PB1) exhibits chromosomes (chr) and cortical ooplasm elements, such as cortical vesicles (cv) and isolated SER tubules (IT). (C) The chromosomes (chr) of the oocyte metaphase II plate appeared slightly displaced towards the upper inner region. Bars: 1  $\mu$ m.

inverted microscope (live oocytes), which confirmed the presence of the first polar body, and in semithin sections (fixed oocytes), which confirmed the presence of the first polar body and the metaphase II plate. Also, in all cases, cytoplasmic integrity was verified through inspection under an inverted microscope and in semithin sections, which confirmed the absence of intracytoplasmic or extracytoplasmic dimorphisms. We started with the donor that provided five oocytes, but TEM observations revealed cytoplasmic immaturity characteristics. Thereafter, we looked at the remaining MII oocytes from the donor to ensure their cytoplasmic maturity. However, all five MII oocytes revealed cytoplasmic immaturity, evidencing an unconformity between nuclear and cytoplasmic

maturity. The current study focuses on the examination of these five oocytes obtained from a single donor.

Previous ultrastructural studies of mature MII oocytes showed the presence of PB1 in the perivitelline space, containing chromosomes and the organelles of the cortical ooplasm (Zamboni *et al.*, 1972; Sathanathan, 1994). An organized metaphase II plate could also be observed in the cortical/subcortical region near the PB1 (Zamboni *et al.*, 1972; Sathanathan, 2013). The oocyte surface presented numerous microvilli (Morimoto, 2009) and CV formed one or more layers next to the oolemma (Sathanathan and Trounson, 1982; Motta *et al.*, 1988), with some cortical regions presenting a higher density of CV than others (Sathanathan and



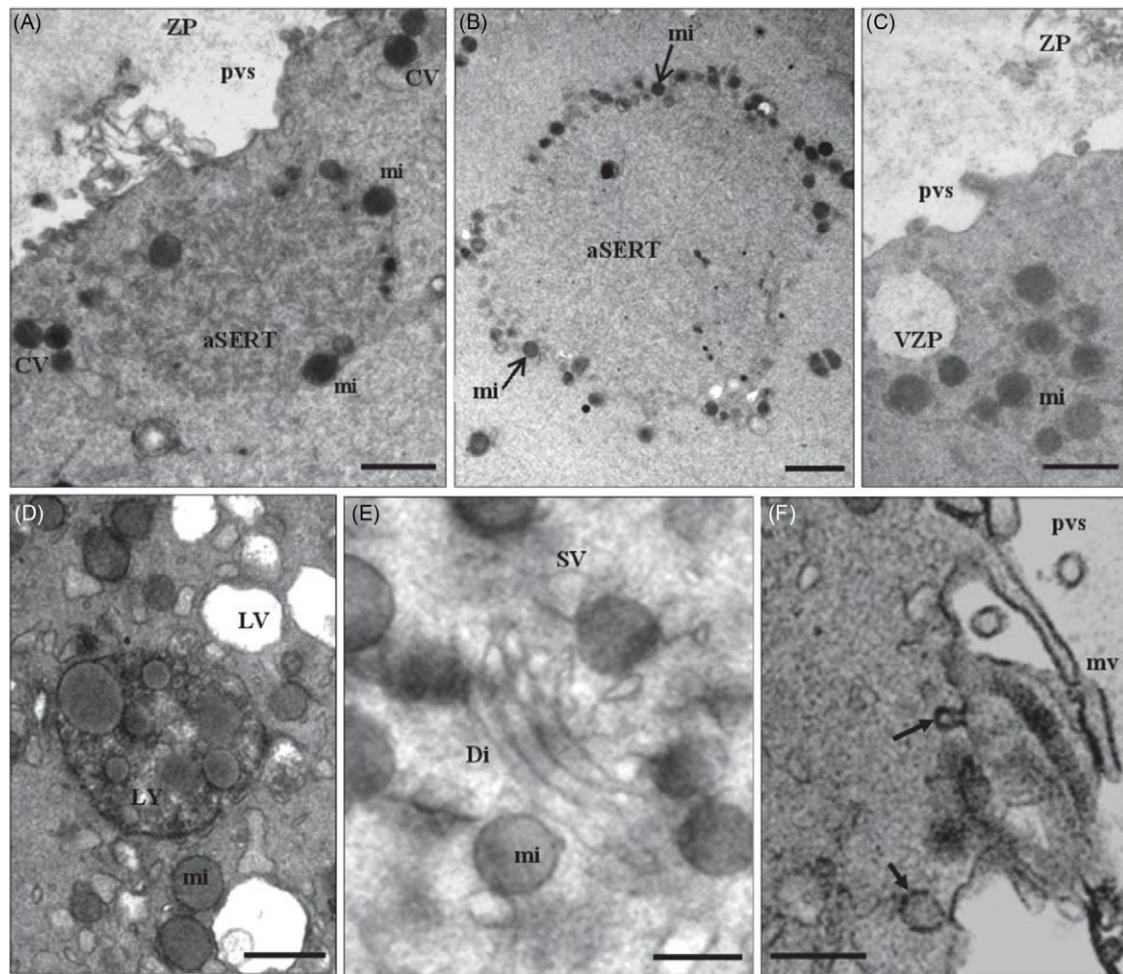
**Figure 3.** Ultrastructural images of metaphase II oocytes at the inner oocyte region. (A) In the inner cytoplasm mitochondria (mi) and smooth endoplasmic reticulum (SER) small (SV), medium (MV) and large (LV) sized vesicles and isolated tubules (IT) can be observed. Note that SER vesicles are barely associated with mitochondria. (B) The inner cytoplasm also presented SER very large vesicles (VLV), with their characteristic bridges (\*).

Trounson, 1982; Sousa and Tesarik, 1994; El-Shafie *et al.*, 2000; Sousa *et al.*, 2016b). The oocyte cortex and subcortex were enriched in SER vesicles of various sizes, SER-IT and aSERT (Sousa and Tesarik, 1994; El-Shafie *et al.*, 2000; Sathananthan, 2013; Sousa *et al.*, 2016b). Mitochondria were described as evenly distributed, surrounding most SER vesicles and aSERT, and forming necklace-like complexes (Sundström *et al.*, 1985; Motta *et al.*, 1988, 2000; El-Shafie *et al.*, 2000; Sathananthan and Trounson, 2000; Morimoto, 2009; Nottola *et al.*, 2014; Sousa *et al.*, 2016b; Trebichalská *et al.*, 2021). Lysosomes were also described (Sathananthan, 2003; Trebichalská *et al.*, 2021), and dictyosomes (Sathananthan *et al.*, 2006; Nottola *et al.*, 2014) and annulate lamella (Sathananthan *et al.*, 2006) were rarely observed. Some organelles and cytoplasmic

structures were not found in mature MII oocytes, such as rough endoplasmic reticulum and multivesicular bodies (Sundström *et al.*, 1985), polyribosomes and lipid droplets (Sathananthan, 1994), just as in donor oocytes (Trebichalská *et al.*, 2021).

We here describe the presence of cortical tiny coated vesicles, which were previously observed in GV and MI oocytes (Pires-Luís *et al.*, 2016; Coelho *et al.*, 2020), and in mature MII oocytes (El-Shafie *et al.*, 2000; Sousa *et al.*, 2016b), indicating exchanges of materials with the exterior by receptor-mediated endocytosis (Hinduja *et al.*, 1990; Pires-Luís *et al.*, 2016; Sousa *et al.*, 2016b; Coelho *et al.*, 2020).

Our present observations confirm those previous findings, but several characteristics of immaturity were also observed in the



**Figure 4.** Ultrastructural images of metaphase II oocytes. (A, B) Aggregates of smooth endoplasmic reticulum tubules (aSERT) in cortical (A) and inner (B) oocyte regions, surrounded by mitochondria (mi). (C) Vesicles containing granular materials (VZP) in the oocyte cortical region. (D) Lysosome (LY) in the inner ooplasm. (E) Dictyosome (Di) in the inner ooplasm. (F) The oocyte surface showed numerous tiny coated vesicles (arrows). ZP: zona pellucida; pvs: perivitelline space; mv: microvilli; SV, LV: small (SV) and large (LV) smooth endoplasmic reticulum vesicles. Bars: (A, C, D) 1  $\mu$ m; (B) 1  $\mu$ m; (E, F) 0.5  $\mu$ m.

ooplasm. These immature characteristics were the following: CV were observed in the oocyte cortex and subcortex, as in GV and MI oocytes (Pires-Luís *et al.*, 2016; Coelho *et al.*, 2020), being that they were expected to be only present in the cortical ooplasm; granular vesicles, suspected of containing ZP-like materials, were observed in the oocyte cortex, as in GV oocytes (Pires-Luís *et al.*, 2016), and although observed in only two oocytes (oocytes 2 and 3), it was expected they would not be found similarly to MI oocytes (Coelho *et al.*, 2020); dictyosomes were observed in the oocyte subcortex, as in GV oocytes (Pires-Luís *et al.*, 2016) and in the IC, as in GV and MI oocytes (Pires-Luís *et al.*, 2016; Coelho *et al.*, 2020), and although observed in only two oocytes (oocytes 1 and 3), they were expected to be not found, although there are two previous reports in mature MII oocytes that observed rare dictyosomes (Sathananthan *et al.*, 2006; Nottola *et al.*, 2014); SER-VLV were observed in the IC in only one oocyte (oocyte 4), but these are a GV characteristic (Pires-Luís *et al.*, 2016); aSERT were missing in oocytes 1–4, and the oocyte exhibiting aSERT (oocyte 5) presented this organelle in all three oocyte regions, as in MI oocytes (Coelho *et al.*, 2020). The aSERT in mature MII oocytes are limited

to the oocyte cortex and subcortex, being critical for the oocyte to develop calcium oscillations at fertilization (Sousa *et al.*, 1996, 1997), without which the oocyte is not activated and no embryo development occurs (Dale *et al.*, 2010).

Therefore, all oocytes exhibited abnormal cortical vesicle positioning in the subcortex and all presented absence or abnormal positioning of aSERT. Additionally, in all cases, mitochondria appeared barely associated with SER-MV and SER-LV, and the metaphase II plate was slightly displaced from the cortical/subcortical region towards the IC. This renders the ooplasm of these donor MII oocytes immature.

By using a quantitative approach, when MII oocytes were compared in their different regions (Table 2), it was shown that there were no significant differences in the Vv of mitochondria among the three oocyte regions, corroborating what was reported in descriptive studies, in which a homogeneous distribution throughout the ooplasm was observed. Although mitochondria were described as the most predominant organelle in the ooplasm (Motta *et al.*, 1988, 2000; Sathananthan and Trounson, 2000; Trebichalská *et al.*, 2021), this was not observed in the present

**Table 1.** Relative volume (Vv) of organelle per total MII oocyte and per MII oocyte regions: cortex, subcortex and inner cytoplasm

Organelle	Vv (organelle, oocyte) mean (%) $\pm$ SEM (Cvar)	Vv (organelle, cortex)	Vv (organelle, subcortex)	Vv (organelle, inner cytoplasm)
Mi	5.49 $\pm$ 0.30 (0.12)	5.66 $\pm$ 0.79 (0.313)	5.79 $\pm$ 0.48 (0.186)	5.33 $\pm$ 0.32 (0.135)
Di	0.01 $\pm$ 0.0085 (1.46)	0	0.01 $\pm$ 0.01 (2.24)	0.018 $\pm$ 0.013 (1.68)
Ly	0.33 $\pm$ 0.11(0.74)	0.03 $\pm$ 0.021 (1.50)	0	0.54 $\pm$ 0.18 (0.739)
CV	0.31 $\pm$ 0.057 (0.403)	1.29 $\pm$ 0.27 (0.462)	0.16 $\pm$ 0.05 (0.30)	0
VZP	0.02 $\pm$ 0.0097 (1.31)	0.084 $\pm$ 0.05 (1.35)	0	0
SV	8.84 $\pm$ 1.87(0.47)	9.67 $\pm$ 1.55 (0.359)	8.74 $\pm$ 1.92 (0.49)	8.40 $\pm$ 1.98 (0.517)
MV	2.01 $\pm$ 0.46 (0.51)	1.04 $\pm$ 0.46 (0.989)	0.74 $\pm$ 0.38 (1.14)	2.75 $\pm$ 0.67 (0.544)
LV	0.79 $\pm$ 0.38 (1.09)	0.12 $\pm$ 0.12 (2.24)	0.012 $\pm$ 0.012 (2.24)	1.27 $\pm$ 0.66 (1.17)
VLV	0.30 $\pm$ 0.29(2,236)	0	0	0.49 $\pm$ 0.49 (2.24)
IT	7.16 $\pm$ 0.69 (0.214)	8.19 $\pm$ 0.83(0.227)	7.22 $\pm$ 1.02 (0.316)	7.2 $\pm$ 0.58 (0.1798)
aSERT	0.61 $\pm$ 0.61 (2.24)	0.98 $\pm$ 0.98(2.24)	0.38 $\pm$ 0.38 (2.24)	0.44 $\pm$ 0.44 (2.24)
Total SER (without VLV)	26.8 $\pm$ 3.31 (0.276)	19.99 $\pm$ 2.17 (0.243)	17.1 $\pm$ 2.71 (0.355)	20.1 $\pm$ 1.46 (0.16)

Results presented as mean, SEM (standard error of the mean: standard deviation/ $n^{1/2}$ ), and CVar (coefficient of variation: standard deviation/mean).

Mi: mitochondria; Di: dictyosomes; Ly: lysosomes; CV: cortical vesicles; VZP: vesicles containing zona pellucida-like material; SER: smooth endoplasmic reticulum; SV: SER small vesicles; MV: SER medium vesicles; LV: SER large vesicles; VLV: SER very large vesicles; IT: SER isolated tubules; aSERT: SER tubular aggregates.

study (Table 2), with quantitative analyses showing that the SER is the most predominant organelle (Table 1) (Mi Vv 5.49% vs total SER Vv 26.8%).

Using strict pairwise comparisons for the three oocyte regions and the Mann–Whitney *U*-test, Bonferroni corrected (Table 2), significant differences were observed for the following and the SER-LV presented a higher prevalence in the IC. This finding contradicts morphological study descriptions, in which lysosomes and SER-LV were seen to be distributed uniformly (Zamboni *et al.*, 1972; Sundström *et al.*, 1985; Motta *et al.*, 1988).

Applying a stereological analysis similar to that previously used in human immature oocytes, GV (Pires-Luís *et al.*, 2016) and MI (Coelho *et al.*, 2020), it was possible to compare quantitatively immature oocytes with the present studied MII oocytes.

Comparing GV with MII oocytes (Table 3), no significant differences were observed in the Vv of mitochondria, lysosomes, CV, VZP and SER-LV. For mitochondria, the Vv was lower in MII oocytes, except in the cortex, and was only significantly lower in the IC; for lysosomes and VZP, no significant differences in the Vv were found along all oocyte regions; in relation to CV, the Vv were similar in the cortex and subcortex; the significant difference observed in the IC were due to the absence of CV in that region of MII oocytes; for SER-LV, the Vv were similar but nevertheless higher in the cortex and IC of MII oocytes, and significantly lower in the subcortex of MII oocytes. Significant differences in all oocyte regions were observed in the Vv of dictyosomes, SER-SV, SER-MV, SER-VLV, SER-IT and total SER. For dictyosomes, the Vv was lower in all regions of MII oocytes; for SER-SV, the Vv was higher in all regions of MII oocytes; for SER-MV, the Vv was higher in all regions of MII oocytes, although not significant in the oocyte subcortex; for SER-VLV, the Vv was lower in all regions of MII oocytes; for SER-IT, the Vv was higher in all regions of MII oocytes; and for total SER ( $P < 0.001$ ), the Vv was higher in all regions of MII oocytes.

Comparing MI with MII oocytes (Table 4), no significant differences in the Vv were found in relation to dictyosomes

(although not significant, the Vv was higher in MII oocytes in the subcortex and IC), lysosomes (except for a significantly lower Vv in the cortex of MII oocytes and a non-significant higher Vv in the IC), CV (although not significant, the Vv was higher in MII oocytes, except for a significantly lower Vv in the IC) and SER-LV (although not significant, the Vv was higher in MII oocytes). Significant differences in the Vv were observed regarding mitochondria, SER-SV, SER-MV, SER-IT, aSERT and total SER. The Vv of mitochondria, SER-IT and aSERT were significantly lower in MII oocytes only when evaluating the total oocyte, with all regions displaying non-significantly lower Vv in MII oocytes. The Vv of total SER was significantly higher when evaluating the total oocyte and IC of MII oocytes, with non-significantly higher Vv in the cortex and subcortex of MII oocytes. The Vv of SER-SV was significantly higher in all regions of MII oocytes. The Vv of SER-MV was significantly lower in the cortex and subcortex of MII oocytes, and not significantly lower in the IC of MII oocytes.

Considering all these quantitative data, a gradual increase in SER elements, as described in morphological studies (Sundström *et al.*, 1985; Nottola *et al.*, 2014) and a decrease in dictyosomes (Sathananthan, 2003; Sathananthan *et al.*, 2006), was noted specifically from GV to MI stages. However, an increase in CV was expected along with oocyte maturation, as well as in aSERT and mitochondria–SER complexes in the oocyte cortex (Nottola *et al.*, 2014), which was not observed. Therefore, the cytoplasm of the oocytes analyzed in this study exhibited both GV and MI features, implying that they contained an immature cytoplasm.

Although a stereological analysis of mature (nuclear and cytoplasmic) MII oocytes is in progress, we think that the presentation of this anomaly (discrepant nuclear and cytoplasmic maturation in MII donor oocytes) deserves anticipated publication due to its immediate clinical importance.

In conclusion, we describe a case of donor MII oocytes exhibiting nuclear maturity simultaneously with cytoplasmic immaturity. If used in a donor programme, these oocytes would not be



**Table 2.** Comparison of the means of organelle relative volume (Vv) between MII oocyte regions: Vv (organelle, cortex), Vv (organelle, subcortex) and Vv (organelle, inner cytoplasm)

Organelle	Kruskal–Wallis test ( <i>P</i> -value*)	Oocyte regions	Mann–Whitney <i>U</i> -test ( <i>P</i> -value*)
Mi	0.677		
Di	0.317		
Ly	0.003	Cortex vs Subcortex (Vv 0.03 vs Vv 0)	0.310
		Cortex vs Inner Cytoplasm (Vv 0.03 vs Vv 0.54)	0.008
		Subcortex vs Inner Cytoplasm (Vv 0 vs Vv 0.54)	0.008
CV	0.002	Cortex vs Subcortex (Vv 1.3 vs Vv 0.16)	0.008
		Cortex vs Inner Cytoplasm (Vv 1.3 vs Vv 0)	0.008
		Subcortex vs Inner Cytoplasm (Vv 0.16 vs Vv 0)	0.008
VZP	0.032	Cortex vs Subcortex (Vv 0.08 vs Vv 0)	0.151
		Cortex vs Inner Cytoplasm (Vv 0.08 vs Vv 0)	0.151
		Subcortex vs Inner Cytoplasm (Vv 0 vs Vv 0)	1.000
SV	0.677		
MV	0.102		
LV	0.009	Cortex vs Subcortex (Vv 0.12 vs Vv 0.01)	1.000
		Cortex vs Inner Cytoplasm (Vv 0.12 vs Vv 1.27)	0.032
		Subcortex vs Inner Cytoplasm (Vv 0.01 vs Vv 1.27)	0.008
VLV	0.368		
IT	0.336		
aSERT	0.980		
Total SER	0.468		

\*Significance set at  $P < 0.05$

Mi: mitochondria; Di: dictyosomes; Ly: lysosomes; CV: cortical vesicles; VZP: vesicles containing zona pellucida-like material; SER: smooth endoplasmic reticulum; SV: SER small vesicles; MV: SER medium vesicles; LV: SER large vesicles; VLV: SER very large vesicles; IT: SER isolated tubules; aSERT: SER tubular aggregates.

Note: Mann–Whitney *U*-test was performed when the Kruskal–Wallis test gave a significant *P*-value.

**Table 3.** Comparison of the means of organelle relative volume (Vv) per oocyte region (organelle, cortex), Vv (organelle, subcortex) and Vv (organelle, inner cytoplasm) between GV oocytes and MII oocytes

Organelle	Total oocyte			Oocyte cortex			Subcortex			Inner cytoplasm		
	Vv			Vv			Vv			Vv		
	GV**	MI	<i>P</i> -value*	GV**	MI	<i>P</i> -value*	GV**	MI	<i>P</i> -value*	GV**	MI	<i>P</i> -value*
Mi	6.3	5.49	1.000	3.6	5.66	0.073	6.0	5.79	0.876	7.2	5.33	0.048
Di	0.5	0.01	<0.001	0.4	0	0.003	0.6	0.01	0.018	0.6	0.018	0.003
Ly	0.2	0.33	0.924	0.1	0.03	0.755	0.1	0	0.268	0.3	0.54	0.149
CV	0.3	0.3	0.485	1.3	1.29	1.00	0.1	0.16	0.343	0.1	0	0.003
VZP	0.05	0.02	0.214	0.2	0.084	0.106	0.02	0	0.432	0	0	0.755
SV	2.5	8.84	<0.001	2.0	9.67	0.003	2.3	8.74	0.005	2.7	8.40	0.040
MV	0.2	2.01	<0.001	0.1	1.04	0.005	0.2	0.74	0.106	0.3	2.75	0.005
LV	0.2	0.79	0.294	0.07	0.12	0.343	0.1	0.012	0.018	0.2	1.27	0.268
VLV	6.9	0.30	<0.001	0.9	0	0.003	1.67	0	0.003	10.1	0.49	0.003
IT	3.2	7.16	<0.001	3.3	8.19	0.018	3.2	7.22	0.010	3.1	7.2	0.010
Total SER (without VLV)	6.1	26.8	<0.001	4.6	19.99	0.003	5.8	17.1	0.003	6.23	20.1	0.003

\*Significance set at  $P < 0.05$ .

\*\*Data are presented as: organelle relative volume (Vv) found in GV oocytes (from Pires-Luis *et al.*, 2016), organelle Vv found in MII oocytes (from Table 1), and the *P*-value obtained after comparison between the GV-organelle Vv and the MII-organelle Vv.

GV: prophase I oocytes (germinal vesicle oocytes); MI: metaphase II oocytes.

Mi: mitochondria; Di: dictyosomes; Ly: lysosomes; CV: cortical vesicles; VZP: vesicles containing zona pellucida-like material; SER: smooth endoplasmic reticulum; SV: SER small vesicles; MV: SER medium vesicles; LV: SER large vesicles; VLV: SER very large vesicles; IT: SER isolated tubules.

Note: aSERT (SER tubular aggregates) not included as not found in GV oocytes.

**Table 4.** Comparison of the means of organelle relative volume (Vv) per oocyte region (organelle, cortex), Vv (organelle, subcortex) and Vv (organelle, inner cytoplasm) between MI oocytes and MII oocytes

Organelle	Oocyte			Cortex			Subcortex			Inner cytoplasm		
	Vv		P-value*	Vv		P-value*	Vv		P-value*	Vv		P-value*
	MI **	MI		MI**	MI		MI**	MI		MI**	MI	
Mi	6.5	5.49	0.023	6.3	5.66	0.690	7.3	5.79	0.095	6.4	5.33	0.421
Di	0.01	0.01	0.713	0.01	0	0.310	0	0.01	0.690	0.01	0.018	1.000
Ly	0.4	0.3	0.074	0.3	0.03	0.032	0.3	0	0.151	0.4	0.54	1.000
CV	0.4	0.3	0.775	0.96	1.29	0.548	0.1	0.16	0.421	0.1	0	0.008
SV	0.2	8.84	<0.001	0.2	9.67	0.008	0.2	8.74	0.008	0.2	8.40	0.008
MV	4.5	2.0	0.001	4.1	1.04	0.032	3.5	0.74	0.016	4.7	2.75	0.421
LV	0.2	0.79	0.935	0.07	0.12	0.310	0	0.012	0.690	0.2	1.27	0.095
IT	9.4	7.2	<0.001	10.4	8.19	0.056	9.6	7.22	0.095	8.8	7.2	0.056
aSERT	0.8	0.3	0.008	1.02	0.98	0.151	1.3	0.38	0.222	0.6	0.44	0.151
Total SER	15.05	26.8	0.013	15.9	19.99	0.151	14.6	17.1	0.548	14.5	20.1	0.032

\*Significance set at  $P < 0.05$ .

\*\*Data are presented as: organelle relative volume (Vv) found in MI oocytes (from Coelho et al., 2020), organelle Vv found in MII oocytes (from Table 1), and the P-value obtained after comparison between the MI-organelle Vv and the MII-organelle Vv.

Mi: mitochondria; Di: dictyosomes; Ly: lysosomes; CV: cortical vesicles; SER: smooth endoplasmic reticulum; SV: SER small vesicles; MV: SER medium vesicles; LV: SER large vesicles; IT: SER isolated tubules; aSERT: SER tubular aggregates.

Note: VZP (vesicles containing zona pellucida-like material) and VLV (SER very large vesicles) not included, as not found in MI oocytes.

fertilized, and therefore this finding is the first demonstration of a possible cause of ART failure in oocyte donor cycles, in which oocyte quality is usually taken for granted.

### Ethics approval and consent to participate

The authors declare that they followed all rules of ethical conduct regarding originality, data processing and analysis, duplicate publication and biological material. All authors critically read the final manuscript and agreed with the author order.

Ethical guidelines were followed conducting the research. According to the National Law on Medically Assisted Procreation (Law 32/2006) and the National Council for Medically Assisted Procreation guidelines (2018), the use of clinical databases and surplus gametes for research may be used under strict individual anonymity and after patient written informed consent. Surplus donor oocytes were used after the patient signed an informed consent agreeing to share their own gamete samples for the present study. The present laboratorial experiments were executed under the Joint Ethics Committee of the Hospital and University, CHUP/ICBAS approval number 2019/CE/P017 (266/CETI/ ICBAS). This work did not involve human or animal experiments and therefore the provisions of the Declaration of Helsinki as revised in Tokyo 2004 do not apply.

**Data availability statement.** The authors confirm that the data supporting the findings of this study is available upon request.

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**Author contributions.** T.S. was responsible for the stereology and statistic work, data analysis, critical discussion, a draft of the manuscript and a critical review of the final manuscript. A.S.P.L. was responsible for study design, stereology and statistic training, electron microscopy image acquisition, data analysis, critical discussion and critical review of the final manuscript. A.A. and E.O. were responsible for oocyte processing to electron microscopy and critical review of the final manuscript. C.L. was responsible for the embryologic work, donor oocyte selection and critical review of the final manuscript. M.F. was

responsible for the psychological evaluation of the donor and the critical review of the final manuscript. E.V.F. and M.B. were responsible for clinical donor selection, ovarian stimulation, oocyte retrieval and critical review of the final manuscript. A.M.C. was responsible for critical review of the final manuscript. R.S. was responsible for critical discussion and review of the final manuscript. M.S. was responsible for study conceptualization, electron microscopy image acquisition, data analysis, critical discussion and writing of the final manuscript. All authors have read and agreed to the order of appearance and the published version of the manuscript.

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**Conflict of Interest.** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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