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## Promoting Extracellular Matrix Remodeling via Ascorbic Acid Enhances the Survival of Primary Ovarian Follicles Encapsulated in Alginate Hydrogels

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### Abstract

The *in vitro* growth of ovarian follicles is an emerging technology for fertility preservation. Various strategies support the culture of secondary and multilayer follicles from various species including mice, non-human primate, and human; however, the culture of early stage (primary and primordial) follicles, which are more abundant in the ovary and survive cryopreservation, has been limited. Hydrogel-encapsulating follicle culture systems that employed feeder cells, such as mouse embryonic fibroblasts (MEFs), stimulated the growth of primary follicles (70–80  $\mu\text{m}$ ); yet, survival was low and smaller follicles (<70  $\mu\text{m}$ ) rapidly lost structure and degenerated. These morphologic changes were associated with a breakdown of the follicular basement membrane; hence, this study investigated ascorbic acid based on its role in extracellular matrix (ECM) deposition/remodeling for other applications. The selection of ascorbic acid was further supported by a microarray analysis that suggested a decrease in mRNA levels of enzymes within the ascorbate pathway between primordial, primary, and secondary follicles. The supplementation of

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### Supporting Information

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ascorbic acid (50 µg/mL) significantly enhanced the survival of primary follicles (<80 µm) cultured in alginate hydrogels, which coincided with improved structural integrity. Follicles developed antral cavities and increased to diameters exceeding 250 µm. Consistent with improved structural integrity, the gene/protein expression of ECM and cell adhesion molecules was significantly changed. This research supports the notion that modifying the culture environment (medium components) can substantially enhance the survival and growth of early stage follicles. *Biotechnol. Bioeng.* 2014;111: 1417–1429.

### Keywords

tissue engineering; regenerative medicine; ovarian follicle development; primary follicle; ascorbic acid; extracellular matrix; biomaterial; alginate hydrogel

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### Introduction

The in vitro growth and maturation of ovarian follicles is an emerging technology with the goal of expanding fertility preservation options for cancer patients (Woodruff, 2007a, b, 2010). Fertility preservation is necessary for girls and women, whose fertility is threatened by lifesaving cancer therapies, such as radiation and chemotherapy, which may diminish the follicle pool and trigger premature ovarian failure. Treatments to restore fertility include the transplantation of cryopreserved ovarian tissue, which may risk the reintroduction of cancer cells (Meirow et al., 2008; Shaw and Trounson, 1997). To avoid this risk, in vitro follicle culture, maturation, and fertilization techniques could potentially be used to produce mature eggs, and subsequently embryos, from cryopreserved ovarian tissue. In vitro culture systems have significantly advanced over the past 35 years (Eppig, 1977; Eppig and O'Brien, 1996; Lenie et al., 2004; Nayudu and Osborn, 1992; O'Brien et al., 2003; Pangas et al., 2003; Torrance et al., 1989); however, further advancements are necessary for the efficient growth of early stage (primordial and primary) follicles and human follicles capable of producing fertilizable eggs.

The three dimensional structure and cell–cell interactions of developing follicles have been maintained by biomaterials, such as alginate hydrogels (Augst et al., 2006; Kreeger et al., 2005, 2006; Pangas et al., 2003). Follicle culture on traditional flat substrates (tissue culture polystyrene) often results in the disruption of the normal follicular architecture (separation of the somatic cells from the oocyte). The maintenance of follicular structure that morphologically resembles in vivo grown follicles has been accomplished by encapsulation within hydrogels. Alginate-based hydrogel culture systems have supported the in vitro growth of secondary and multilayer mouse, non-human primate, and human follicles (Shikanov et al., 2009; West et al., 2007; Xu et al., 2006b, 2009a,b, 2011, 2013). Mouse follicles have produced eggs capable of fertilization and live births (Xu et al., 2006a). Furthermore, these systems provide a controlled and modifiable environment in which to study the mechanisms governing follicle development (Hornick et al., 2012; Kreeger et al., 2005, 2006; Parrish et al., 2011; Skory et al., 2013; West et al., 2007; Xu et al., 2006b).

As culture systems continue to advance, the focus is shifting to early stage (primary and primordial) follicles, which are more abundant in the ovary and survive cryopreservation,

yet the biology is not well understood and successful culture has been limited (Abir et al., 2006; Itoh and Hoshi, 2000; Lenie et al., 2004; Muruvi et al., 2005, 2009; Saha et al., 2000). Early stage follicles have been grown via in situ organ culture or two-step culture systems (Eppig and O'Brien, 1996; Jin et al., 2010), in which ovarian tissue fragments containing early stage follicles are cultured in vitro. The tissue and stromal cells within the ovarian tissue provide physical support as well as signals that stimulate follicle growth. In alginate-based culture systems, the growth of isolated primary follicles has been achieved via co-culture with ovarian stromal cells (theca-interstitial cells; Tingen et al., 2011), mouse embryonic fibroblasts (MEFs; Tagler et al., 2012), and multiple follicles (Hornick et al., 2013). The hydrogel physically supports follicle structure (Hornick et al., 2012), while the feeder cells secrete paracrine factors that stimulate growth. These co-culture methods have been effective for early secondary follicles (90–100  $\mu\text{m}$ ), but primary follicles (70–80  $\mu\text{m}$ ) have low survival rates (Tagler et al., 2012). Therefore, these co-culture systems require additional factors to improve the survival of growing follicles and potentially culture smaller follicles.

In this report, we aimed to improve the survival and growth of primary follicles (60–70  $\mu\text{m}$ ) encapsulated in alginate hydrogels and co-cultured with MEFs (Tagler et al., 2013). The limited growth of early stage follicles results from a loss of follicle structure, which coincided with a breakdown of the follicular basement membrane that may result from damage during follicle isolation. Ascorbic acid, which has been associated with extracellular matrix (ECM) deposition/remodeling, has been previously identified to improve secondary and multilayer follicle (150–200  $\mu\text{m}$ ) survival (Murray et al., 2001; Rose et al., 1999; Thomas et al., 2001); and herein, we investigated ascorbic acid as a means to improve primary follicle culture in alginate hydrogels. In addition to investigating the influence of ascorbic acid on the survival and growth of primary follicles, we explored the underlying mechanisms, such as oxidative stress (antioxidant) and ECM gene/protein expression.

## Materials and Methods

### Animals and Materials

Follicles were isolated from the ovaries of CD1 female mice. Mice were maintained in accordance with the policies of the National Institutes of Health and Northwestern University's Animal Care and Use Committee. A temperature, humidity, and light (12 hL/12 hD) controlled barrier facility within Northwestern University's Center of Comparative Medicine (Chicago, IL) was used to house and breed the mice. Mice were provided with food and water ad libitum. The mice were fed Teklad Global (Madison, WI) irradiated chow (2,919 or 2,916) that does not contain soybean or alfalfa meal but does contain minimal phytoestrogens. Unless otherwise specified, all chemical were purchased from Sigma–Aldrich (St Louis, MO) and medium formulations were purchased from Life Technologies (Calsbad, CA). Sodium alginate (55–65% guluronic acid) was provided by FMC BioPolymers (Philadelphia, PA).

## Gene Expression During Early Stage Follicle Development (mRNA Microarray)

Previously established procedures (Xu et al., 2006b) for ovary/follicle isolation were followed with slight modifications. Primordial follicles were isolated from the ovaries of mice at post-natal day 3 and day 4 (Skory et al., 2013). Primary follicles (70–90  $\mu\text{m}$  in diameter) were mechanically isolated from day 10 ovaries. Two-layered secondary follicles (100–130  $\mu\text{m}$  in diameter) were mechanically isolated from day 12 ovaries. Follicles were then aspirated. Each sample was collected in triplicate. RNA isolation from the ovary (primordial follicles) or from the entire follicles (primary and two-layered secondary) was performed as previously described (Skory et al., 2013). RNA was hybridized in MouseRef-8 v2 Expression BeadChips from Illumina and time series transcriptomics were employed to determine the genes that were differentially expressed over time. Normalization, transformation, and determination of adequate and present probes were performed as previously described (Skory et al., 2013). Briefly, data were transformed and normalized with the variance stabilization transformation method (Lin et al., 2008) and normalized by robust spline normalization (Du et al., 2008). Differentially expressed genes over time were assessed using limma (Smyth, 2004). Differentially expressed genes with a fold change per day of greater than 1.4 and FDR (false discovery rate) corrected *P*-value less than 0.01 were considered significant.

## Follicle Isolation and Encapsulation

Previously established procedures (Tagler et al., 2012) for follicle isolation and alginate encapsulation were followed with slight modifications. Briefly, individual secondary (diameter: 90–100  $\mu\text{m}$ ) and primary follicles (diameter: 60–80  $\mu\text{m}$ ) were mechanically isolated from 7–10-day-old CD1 mice using insulin gauge needles in dissection medium (Leibovitz's L-15 medium supplemented with 1% fetal bovine serum and 0.5% penicillin–streptomycin). Follicles were separated into size classes based on average initial diameter (60, 70, 80, 90, and 100  $\mu\text{m}$ ). After isolation, the follicles were washed and stored in maintenance medium (Minimum Essential Medium Eagle Alpha Modification [aMEM] supplemented with 1% fetal bovine serum and 0.5% penicillin–streptomycin). Follicles were then transferred into 5–6  $\mu\text{L}$  solutions of 0.25% alginate and immersed into a 50 mM  $\text{CaCl}_2$  and 140 mM NaCl solution for 1–2 min to crosslink. Encapsulated follicles were then co-cultured with mouse embryonic fibroblasts (MEFs).

## Mouse Embryonic Fibroblast (MEF) Co-Culture

Follicles were co-cultured with MEFs according to previously established procedures (Tagler et al., 2012). Briefly, Inactivated MEFs (Life Technologies #S1520-100) were thawed, resuspended, and seeded according to the manufacturer's procedures. A hemocytometer was used to determine cell concentration, and viability was determined to be approximately 90% via trypan blue staining. MEFs were seeded at  $2.0 \times 10^4$  viable cells/well in a 96-well flat-bottom culture plate (Corning Costar #3596). After 20 h of culture (overnight), the cells were washed once in 100  $\mu\text{L}$  of PBS for 5 min and 100  $\mu\text{L}$  of fresh follicle growth medium was added to each well. Following follicle isolation and encapsulation, individual alginate beads were placed into each well. The co-culture was

conducted at 37°C in 5% CO<sub>2</sub> for 6–14 days. Every 2 days, half the medium (50 µL) was replaced with fresh growth medium.

### **Ascorbic Acid (AA) and Control Growth Medium Formulations**

The ascorbic acid (AA+) growth medium consisted of αMEM medium supplemented with 3 mg/mL bovine serum albumin (BSA; MP Biomedicals, Solon, OH), 1 mg/mL bovine fetuin, 5 µg/mL insulin, 5 µg/mL transferrin, 5 ng/mL selenium (ITS), 10 mIU/mL recombinant human follicle-stimulating hormone (FSH) (A.F. Parlow, NHPP, NIDDK), and 50 µg/mL sodium l-ascorbic acid (Sigma #A4034; Thomas et al., 2001). Fresh AA growth medium was made every 2 days. The control growth medium without ascorbic acid (–AA) consisted of αMEM supplemented with BSA, fetuin, ITS, and FSH.

### **Follicle Survival and Growth Measurements**

Images of follicles were collected every other day using an inverted Leica DM light microscope (Leica, Wetzlar, Germany). Follicle diameters were measured using ImageJ software (National Institutes of Health, Bethesda, MD). An average of two perpendicular diameter measurements was used for each follicle at each time point. Survival was determined via follicle/oocyte morphology. Non-growing follicles or follicles with fragmented, shrunken, degenerated, or detached oocytes were considered dead.

### **Confocal Imaging**

Follicles were fixed in 4% paraformaldehyde for 45 min at room temperature and then washed in a blocking solution containing 0.2% sodium azide, 2% goat serum, 1% bovine serum albumin, and 0.1% Triton X-100. Next, follicles were stained with primary antibodies for laminin (1:100, rabbit anti-mouse, Sigma #L9393), rhodamine phalloidin (f-actin; 1:50, Life Technologies #R415), and DAPI (4',6-diamidino-2-phenylindole, dihydrochloride) (1 µg/mL, Life Technologies #D1306). Follicles were then washed three times for 10 min in wash buffer and then incubated at room temperature for 1 h with the secondary antibody (Alexa Fluor® 568 goat anti-rabbit IgG, 1:50, Life Technologies #A-11011). Finally, follicles were mounted on Teflon printed microscope slides (Electron Microscopy Sciences, PA). Samples were imaged on Zeiss LSM 510 Meta Confocal Microscope using 40× oil objectives.

### **Oxidative Stress**

The OxiSelect™ In Vitro ROS/RNS Assay Kit (Cell Biolabs #STA-347) was used to test conditioned medium for reaction oxygen and nitrogen species (ROS/RNS). Half the growth medium (50 µL) was collected on days 2, 4, and 6 and stored in –80°C. The protocol provided by the manufacturer was followed.

### **Extracellular Matrix (ECM) and Cell Adhesion Molecules Gene Expression (qRT-PCR)**

The mouse extracellular matrix (ECM) and cell adhesion molecules qRT-PCR (Quantitative Real-Time PCR) array (Qiagen, Hilden, Germany) was utilized to determine differences in the gene expression (mRNA levels) between follicles (<75 µm) cultured in vitro in the presence and absence of ascorbic acid. RNA extraction, cDNA synthesis, and qRT-PCR

array were performed according to the manufacturer's instructions. Follicles in each group were removed from alginate beads on day 6 of culture by incubating in 10 mIU/mL alginate lyase for 10 min. The follicles were then flash frozen in liquid nitrogen in extraction buffer and stored at  $-80^{\circ}\text{C}$ . RNA was extracted using RNeasy Micro Kit (Qiagen) and cDNA was generated using pathway specific primers to enrich for mouse ECM and cell adhesion molecules genes. Genomic DNA was also eliminated in this step. Amplified cDNA was added to RT<sup>2</sup> SYBR Green qPCR master mix (Qiagen) and 10  $\mu\text{L}$  was added to a 384-well plate, with each well containing pre-dispensed gene-specific primer sets. Included on the plate was 1 genomic DNA control, 3 reverse transcription controls (RTC), and 3 positive PCR controls (PPC). The plate was sealed and loaded onto an ABI 7900 HT thermal cycler. ABI 7900 HT SDS software version 2.3 (Applied Biosystems, Foster City, CA) was used to determine the threshold cycle ( $C_t$ ) values for all genes on the qPCR array. Using the RT<sup>2</sup> Profiler PCR Array Data Analysis software version 3.5 (Qiagen), the  $C_t$  values were normalized against the average  $C_t$  of housekeeping genes (Actb, B2m, and Gusb) to generate delta  $C_t$  ( $\Delta C_t$ ). The  $2^{-\Delta C_t}$  method (Livak and Schmittgen, 2001) was used to determine fold change. Fold-change ( $2^{-\Delta C_t}$ ) was calculated as the normalized gene expression ( $2^{-\Delta C_t}$ ) of follicles cultured in the presence of ascorbic acid (AA+) divided by the normalized gene expression ( $2^{-\Delta C_t}$ ) of follicles cultured in the absence of ascorbic acid (AA-).

### Western Blot (Immunoblot) Analysis and Protein Quantification

Follicles ( $<75 \mu\text{m}$ ) were cultured in growth medium with or without ascorbic acid for 7 days. Surviving follicles were removed from alginate beads by incubating with 50  $\mu\text{L}$  of  $\alpha\text{MEM}$  containing 10 IU/mL alginate lyase for 10 min. The follicles were then lysed in lysis buffer (20 mM Tris-HCl, pH 8.0, 137 mM NaCl, 10% glycerol, 1% NP-40, 2 mM EDTA) and flash frozen in liquid nitrogen by treatment group. Samples were stored in  $-80^{\circ}\text{C}$  until further analysis. After thawing, the lysates were electrophoresed on 4–12% SDS-PAGE gels and then transferred to nitrocellulose for immunoblot analysis. The blots were probed by polyclonal anti-fibronectin antibody (abcam, Cambridge, MA) overnight in  $4^{\circ}\text{C}$  followed by anti-rabbit secondary antibody conjugated to horseradish peroxidase (Zymed, San Francisco, CA). Proteins were detected by ECL primer (GE HealthCare Life Sciences, Pittsburgh, PA) and exposed to X-ray film (Kodak, Rochester, NY). The same blot was stripped using stripping buffer (Thermo Scientific) and re-probed with monoclonal anti- $\alpha$ -tubulin (Sigma, St. Louis, MO) followed by an anti-mouse secondary antibody conjugated to horseradish peroxidase. The blot was detected by the same method described above. Protein quantification was done using densitometric analysis on ImageJ Imaging Software.

### Statistical Analysis

Secondary and primary follicle survival and growth (in the presence and absence of ascorbic acid) data were collected via ten independent cultures of approximately 40–80 follicles each. The extracellular matrix (ECM) and cell adhesion protein gene expression (mRNA) (qRT-PCR) data were collected via four biological replicates consisting of approximately 30 follicles per condition. Western blot data were collected via three independent cultures of approximately 20–50 follicles each. Oxidative stress measurements were collected via 10–12 independent conditioned medium samples. Statistical analysis was performed using Microsoft Excel and GraphPad Prism. Follicle growth and oxidative stress data were



analyzed using one-way analysis of variance (ANOVA) followed by Tukey for single time point comparisons. Survival data were analyzed using the Ka-plan–Meier log-rank test. For gene expression, *P*-values were calculated based on the Student's *t*-test of the replicate  $2^{-Ct}$  values for each gene in the control and ascorbic acid treated groups. Western blot data were analyzed using a one-tailed unpaired Student's *t*-test. A *P*-value less than 0.05 was considered statistically significant.

## RESULTS

### Primary Follicle Survival With Mouse Embryonic Fibroblast (MEF) Co-Culture

Primary follicles (average initial diameter of 60–70  $\mu\text{m}$ ) co-cultured with MEFs had low survival rates and exhibited the loss of follicle structure (Fig. 1). The survival rates were 23% for 70  $\mu\text{m}$  (day 14), and 0% for 60  $\mu\text{m}$  follicles (day 18). In non-surviving follicles, the oocyte typically separated from the somatic cells and degenerated. Analysis of the degenerating follicles by confocal imaging for laminin demonstrated the disruption of the basement membrane, despite the presence of the encapsulating alginate hydrogel. This phenotype suggested that the extracellular matrix (ECM) and/or basement membrane were insufficient to support follicle development.

Literature searches identified ascorbic acid as a factor that influences ECM deposition/remodeling (Murray et al., 2001; Thomas et al., 2001), and microarray data was utilized to determine if the expression of genes within the ascorbate pathway changed during the primordial-to-primary and primary-to-secondary follicle transitions (Supplemental Table I). Follicles at the primordial and primary stage had a significant decline in mRNA levels for genes that encode enzymes within with the ascorbate/aldarate pathway, and a similar decline was observed in the primary-to-secondary follicle transition. This decline in expression of enzymes suggests that endogenous production of ascorbic acid may decline *in vivo*, as ascorbic acid may not be required given the presence of stromal cells and ECM. However, a decline in ascorbic acid production may limit the ability of the follicle to stimulate ECM deposition in response to damage that occurred due to isolation. The combination of basement membrane breakdown and altered expression of enzymes within the ascorbate pathway led to the hypothesis that ascorbic acid could limit basement membrane breakdown during early stage follicle culture. Hence, this hypothesis was investigated via ascorbic acid supplementation as a means to improve the structural integrity of developing primary follicles.

### Follicle Survival in the Presence ( $\pm$ AA) of Ascorbic Acid

Secondary (average initial diameter of 90–100  $\mu\text{m}$ ) and primary follicles (average initial diameter of 60–80  $\mu\text{m}$ ) were cultured in the presence of 50  $\mu\text{g}/\text{mL}$  ascorbic acid for 14–18 days. Ascorbic acid significantly enhanced the survival of 60–70  $\mu\text{m}$  primary follicles (Fig. 2, and Table I) encapsulated in alginate hydrogels and co-cultured with MEFs, and these follicles cultured demonstrated similar morphologies to *in vivo* grown follicles (Fig. 3). In the presence of ascorbic acid, the survival rates at the end of follicle culture were 48% for 70  $\mu\text{m}$  (day 14), and 17% for 60  $\mu\text{m}$  follicles (day 18), a significant increase relative to the absence of ascorbic acid. The survival of 80, 90, and 100  $\mu\text{m}$  follicles was not significantly

impacted by the presence of ascorbic acid (Supplemental Fig. 1). Moreover, for all size classes, no significant differences were observed with respect to follicle growth between conditions.

### Oxidative Stress

Ascorbic acid was initially proposed based on its effects on ECM deposition/remodeling; however, ascorbic acid is also an antioxidant and we subsequently investigated whether the antioxidant properties contributed to follicle survival (Luck et al., 1995; Meister, 1994). Oxidative stress was measured as the total reactive oxygen species (ROS) and reactive nitrogen species (RNS) in the conditioned medium (Fig. 4). The control medium (-AA) had ROS and RNS levels that did not significantly change throughout culture. Follicles cultured in the presence of ascorbic acid +AA had similar levels of oxidative stress at day 2, with modest increases at day 4 and 6. As a control, an alternative antioxidant, glutathione (GSH), was added to the culture medium. Supplementing glutathione (0.2, 1, and 2 mM) did not improve follicle survival or growth compared to the control (-AA and -GSH; Supplemental Table II). Measurement of ROS and RNS levels with 0.2 mM glutathione indicated that the oxidative stress was comparable to the ascorbic acid condition (Fig. 4). The greatest concentration of glutathione tested (2 mM) substantially increased the ROS and RNS levels at the initial time point, which declined with time in culture. The modest levels of ROS and RNS in the control culture media combined with the inability of glutathione to improve follicle survival suggested that the antioxidant activity of ascorbic acid was not the critical factor influencing primary follicle survival.

### Extracellular Matrix (ECM) and Cell Adhesion Molecules Gene (mRNA) Expression

The previously reported link between ascorbic acid and ECM deposition/remodeling (changes in protease and protease inhibitor expression; Murray et al., 2001; Thomas et al., 2001) was further investigated via the gene expression of 84 ECM and cell adhesion molecules. Primary follicles (<75  $\mu$ m) were cultured for 6 days in the absence (-AA) and presence (+AA) of ascorbic acid. Significantly different gene expression levels between conditions were identified (Fig. 5, Table II [selected data], and Supplemental Table III [all data]). Of the 84 genes tested, 22 significantly changed between conditions ( $P < 0.05$ ). Only seven of the significantly different genes had a fold change greater than two. Matrix metalloproteinase 10/12 (MMP-10/12), fibronectin 1 (Fn1), and collagen type IV alpha 2 (Col4a2) had a greater than twofold increase in gene expression. Cadherin 1 (Cdh1), ectonucleoside triphosphate diphosphohydrolase 1 (Entpd1), and integrin alpha 2 (Itga2) had a greater than twofold decrease in gene expression. Col4a2, Fn1, and Entpd1 are basement membrane components, and MMP-10/12 are proteases that can degrade collagen and fibronectin. Cdh1 and Itga2 are cell-adhesion molecules.

### Western Blots

The variations in gene expression were confirmed by performing Western blots for fibronectin, which exhibited significantly up-regulated gene expression in the presence of ascorbic acid. Primary follicles (<75  $\mu$ m) were cultured for 7 days. Follicles cultured with ascorbic acid (+AA) produced a denser band for fibronectin than those cultured without



ascorbic acid (–AA; Fig. 6). After determining the optical densities of the bands, normalized values were determined by calculating the ratio of the density of the band for fibronectin to that of  $\alpha$ -tubulin for each given sample. The +AA group was significantly greater than the –AA group.

## Discussion

The in vitro culture of primordial and primary follicles has been challenging (Abir et al., 1999; Itoh and Hoshi, 2000; Muruvi et al., 2005, 2009; Picton et al., 2008; Saha et al., 2000; Xu et al., 2013), in part due to the inability to maintain follicle structure. Follicle culture systems that utilize flat surfaces (Lenie et al., 2004) can support the growth of mouse primary follicles (80–90  $\mu\text{m}$ ), although physical manipulations are required to prevent somatic cell attachment and migration away from the oocyte. Some success has been achieved via in situ organ culture or two-step culture systems (Eppig and O'Brien, 1996; Jin et al., 2010). As with organ culture, hydrogels that encapsulate follicles can maintain the architecture and preserve cell–cell connections (Hornick et al., 2012). However, for the culture of early stage follicles in hydrogels, co-culture with supporting cells, such as stromal cells, is necessary to promote survival and growth (Hornick et al., 2013; Tagler et al., 2012; Tingen et al., 2011). More recently, mouse embryonic fibroblast (MEF) co-culture supported the growth of early secondary (90–100  $\mu\text{m}$ ) and late primary (70–80  $\mu\text{m}$ ) follicles; however, the survival of developing primary follicles was low and smaller follicles (<70  $\mu\text{m}$ ) rapidly degenerated (Tagler et al., 2012). To improve the survival of primary follicles, this study investigated ascorbic acid in order to enhance the extracellular matrix (ECM) and prevent separation of the oocyte from the somatic cells.

Ascorbic acid enhanced the survival of primary follicles (60–70  $\mu\text{m}$ ), complementing previous research that observed the improved survival of secondary and multilayer follicles (150–200  $\mu\text{m}$ ; Murray et al., 2001; Rose et al., 1999; Thomas et al., 2001). Ascorbic acid has been used as a component of various follicle/oocyte culture systems and studies (Dalvit et al., 2005; Durlinger et al., 2001; Eppig et al., 1998; Kim et al., 2004; Mao et al., 2002; McLaughlin and Telfer, 2010; Nation and Selwood, 2009; Nayudu et al., 2003; Nilsson et al., 2002; O'Brien et al., 2003; Rossetto et al., 2009; Tao et al., 2010; Telfer et al., 2008; Vitt et al., 1998; Wu et al., 2001). The study herein focused on primary follicles, and the presence of ascorbic acid improved follicle survival. Non-surviving follicles were observed to extrude the oocyte from the somatic cells, an obvious disruption in the basement membrane of the follicle. In the presence of ascorbic acid, follicle survival increased, and it was hypothesized that enhanced ECM would maintain the follicle architecture. The basement membrane, and more generally the ECM within the follicle, plays a critical role in various ovarian cell functions, such as adhesion, migration, survival, differentiation, and proliferation, via mechanical and chemical signals (Berkholtz et al., 2006b; Irving-Rodgers and Rodgers, 2006; Kreeger et al., 2006; Rodgers et al., 2003; Woodruff and Shea, 2007). The composition of the ECM in the murine follicles has been characterized, with collagen type IV and fibronectin present in the theca cell layer and basement membrane throughout follicle development (Berkholtz et al., 2006a; Rodgers et al., 1998).

The hypothesized connection between ascorbic acid and ECM deposition/remodeling in primary follicles was investigated through gene expression for numerous ECM proteins or enzymes. Collagen type IV, fibronectin, matrix metalloproteinase 10 (MMP-10), and matrix metalloproteinase 12 (MMP-12) were significantly different in the presence of ascorbic acid. Consistent with our results, previous studies investigating ascorbic acid identified significant changes in collagen (Hulmes, 1992; Luck et al., 1995; Ono et al., 1990; Pinnell, 1985). The enzymes MMP-10 and MMP-12 are reported to contribute to remodeling of collagen type IV, fibronectin, elastin, and plasminogen (Smith et al., 2002). Moreover, significant changes have been observed in matrix metalloproteinase (MMP-9; Thomas et al., 2001) and tissue inhibitor of metalloproteinase 1 (TIMP-1; Murray et al., 2001) due to ascorbic acid. Interestingly, changes in MMP-9 and TIMP-1 were not observed in our study. These differences may be attributable to follicle size class (primary vs. secondary/multilayer), animal species (mouse vs. bovine), or culture system (alginate hydrogel vs. tissue culture plates). The proposed hypothesis is that changes in gene expression function to strengthen the basement membrane of the developing primary follicles, preventing the loss of follicular structure (oocyte separating from the somatic cells). In contrast to secondary follicles, which have been cultured effectively without ascorbic acid, primary follicles may lack the ECM required to maintain follicular structure.

In addition to ECM proteins, significant changes in the gene expression of cell adhesion molecules were observed. Cadherin 1, which has been associated with oocyte/follicle development (Blaschuk and Farookhi, 1989; Carabatsos et al., 2000; Machell et al., 2000; Mackay et al., 1999; Wang and Roy, 2010), was significantly down-regulated by ascorbic acid. Integrin alpha 2, which was similarly downregulated with ascorbic acid, has been observed on granulosa cells boarding the basement membrane (Giebel and Rune, 1997; Giebel et al., 1996; Wehrenberg and Rune, 2000; Yamada et al., 1999). The down-regulation of both cadherin 1 and integrin alpha 2 is consistent with previous research that observed decreasing expression during primordial-to-primary-to-secondary follicle development (Giebel et al., 1996; Machell et al., 2000). These results demonstrate that ascorbic acid stimulates changes in cell adhesion molecules that are consistent with normal follicle development.

Ascorbic acid may also function as an antioxidant or free radical scavenger to improve follicle survival (Luck et al., 1995; Meister, 1994). Oxidative stress or the imbalance of reactive oxygen species (ROS) and/or reactive nitrogen species (RNS) is associated with cell damage/death and various reproductive diseases (Agarwal et al., 2012). However, in this study, ascorbic acid was not observed to lower the levels of ROS and RNS in conditioned medium from primary follicles co-cultured with MEFs. Previous studies have connected ascorbic acid to granulosa and theca cell death/proliferation (Duleba et al., 2004; Thomas et al., 2001; Tilly and Tilly, 1995). Other antioxidants, such as glutathione and vitamin E, also have been observed to improve embryo/follicle/ovary culture (Eppig et al., 2000; Nugent et al., 1998; Olson and Seidel, 2000; Tao et al., 2010; Wang et al., 2002). In this study, glutathione was tested but did not improve primary follicle survival, which suggested that antioxidant activity was not the critical factor. Stage-specific differences between secondary/multilayer follicles from previous studies and primary follicles reported herein may explain

this discrepancy. Alternatively, glutathione may not be deficient or a limiting factor in our system.

In conclusion, we have demonstrated that ascorbic acid improves the survival of primary mouse ovarian follicles (60–70  $\mu\text{m}$ ). This culture system combined biomaterials (alginate hydrogels), co-culture with MEFs, and medium containing ascorbic acid to simultaneously support follicle structure/survival and stimulate growth. Without ascorbic acid, 60  $\mu\text{m}$  primary follicles rapidly lost structure and degenerated within 10 days. In contrast, follicles cultured in the presence of ascorbic acid survived for 18 days and increased to diameters exceeding 250  $\mu\text{m}$ . The mechanism of action involves gene/protein expression changes associated with ECM and cell adhesion molecules, though additional effects are possible. Future research may lead to novel strategies for in vitro follicle culture and fertility preservation.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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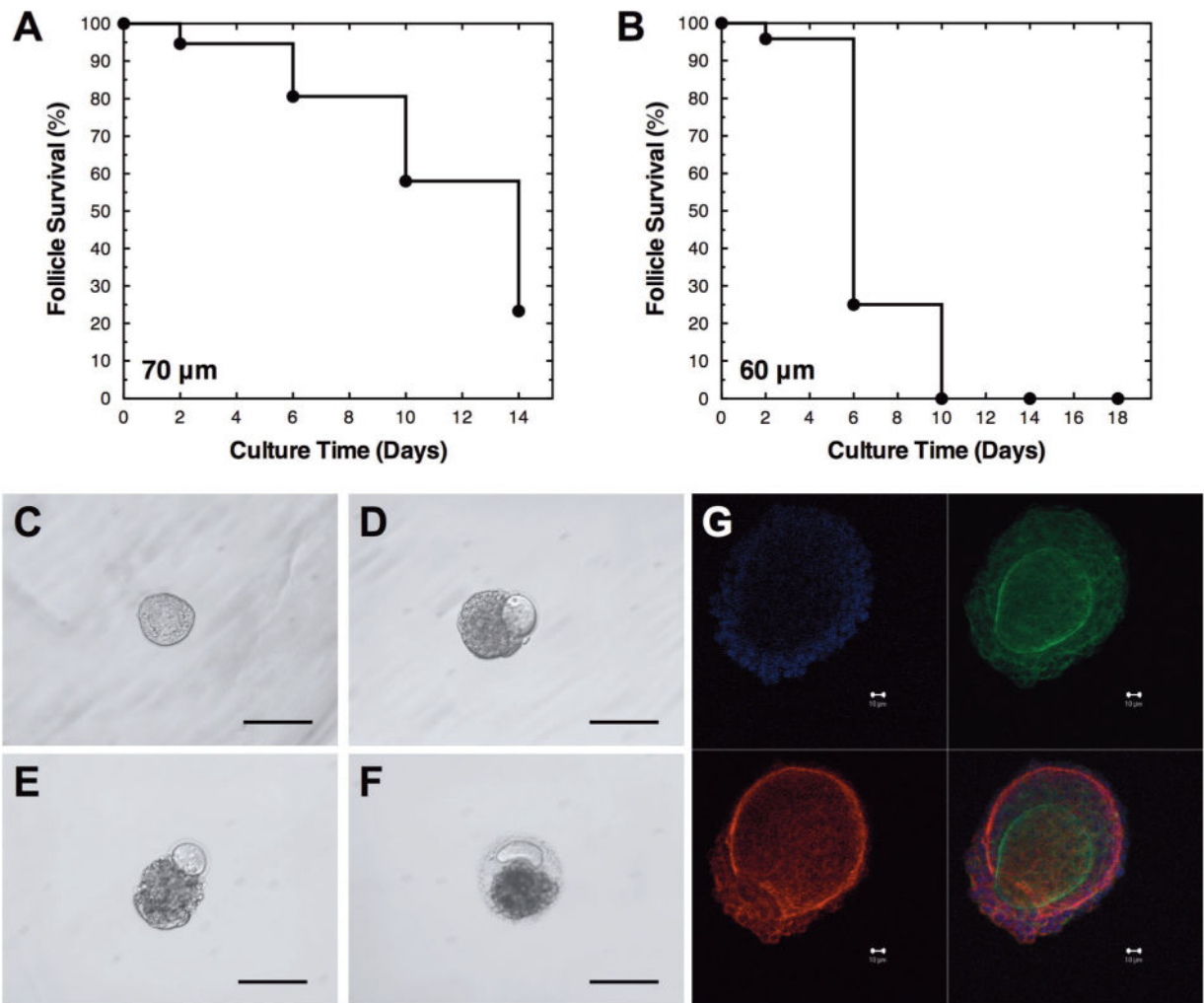
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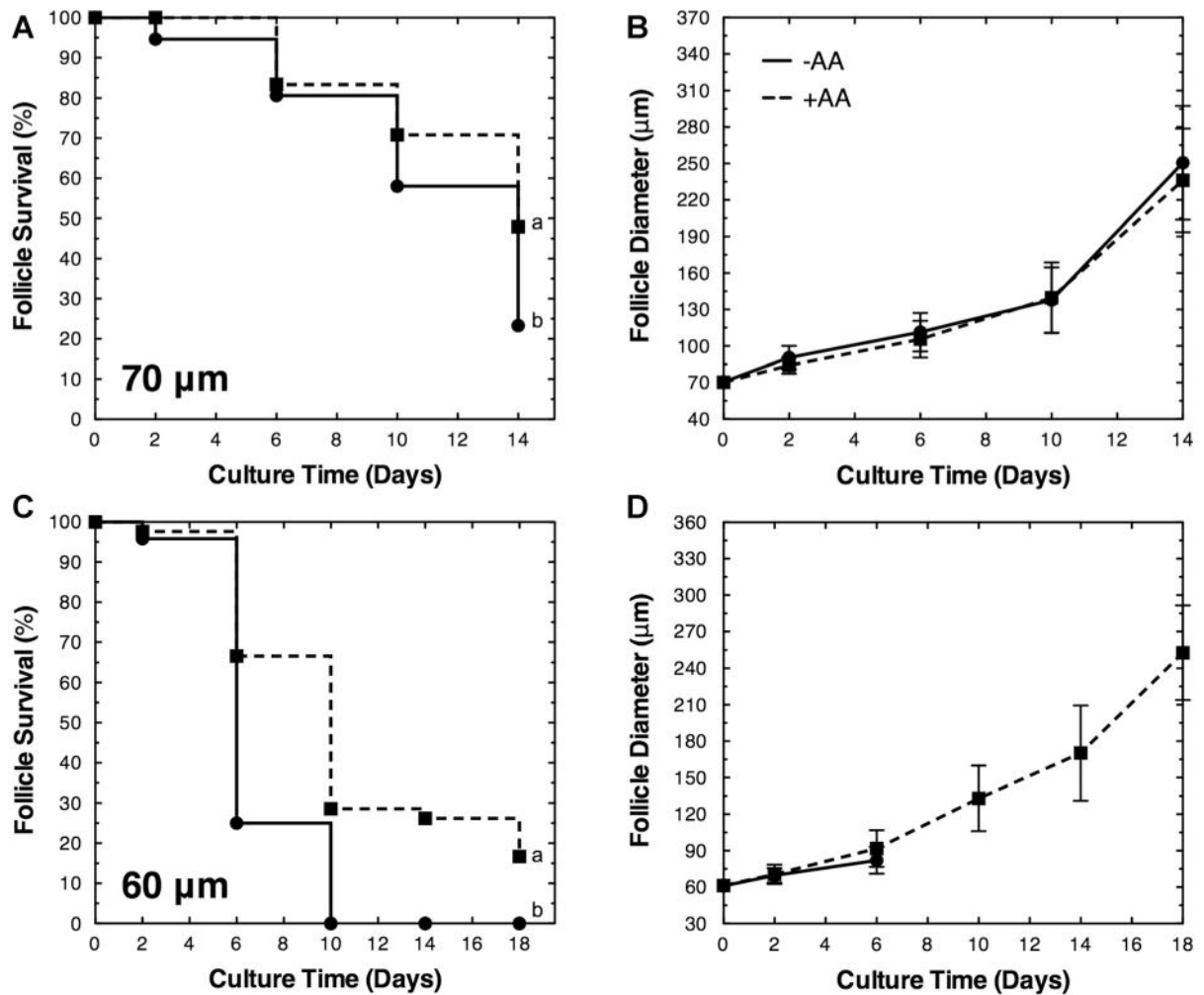
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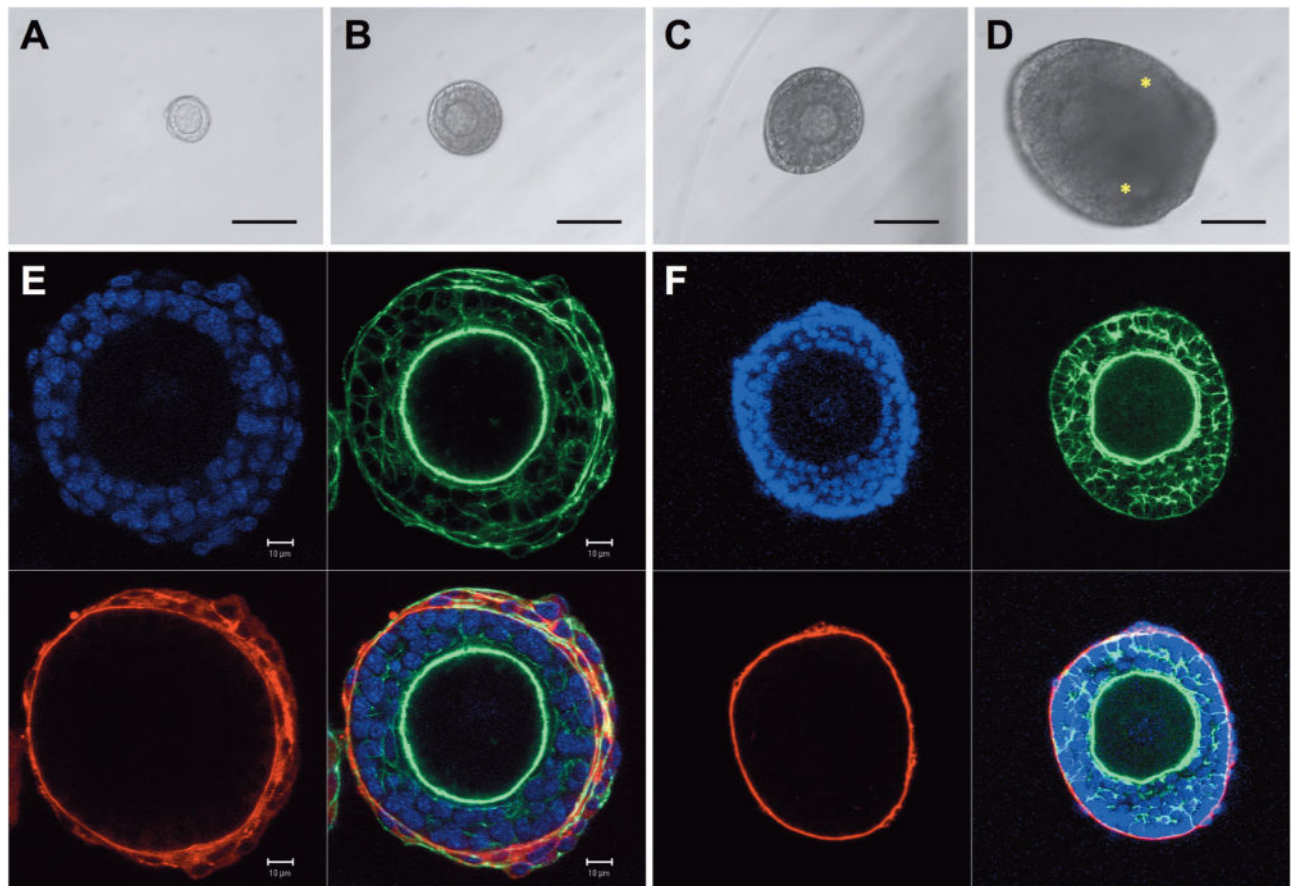


**Figure 1.**

Primary follicle survival with mouse embryonic fibroblast (MEF) co-culture. The survival of (A) 70 μm and (B) 60 μm primary follicles cultured with MEFs. Representative images of primary follicles on (C) day 0, (D) day 6, (E) day 10, and (F) day 16. (G) Confocal images of representative non-growing follicle cultured on day 6 (red, laminin; green, f-actin; and blue, nucleus). Follicles have low survival, which is typically characterized by the loss of follicle structure, extruded oocyte, ruptured basement membrane, and degeneration. The scale bars represent (C–F) 100 μm and (G) 10 μm. Sample sizes are presented in Table I.

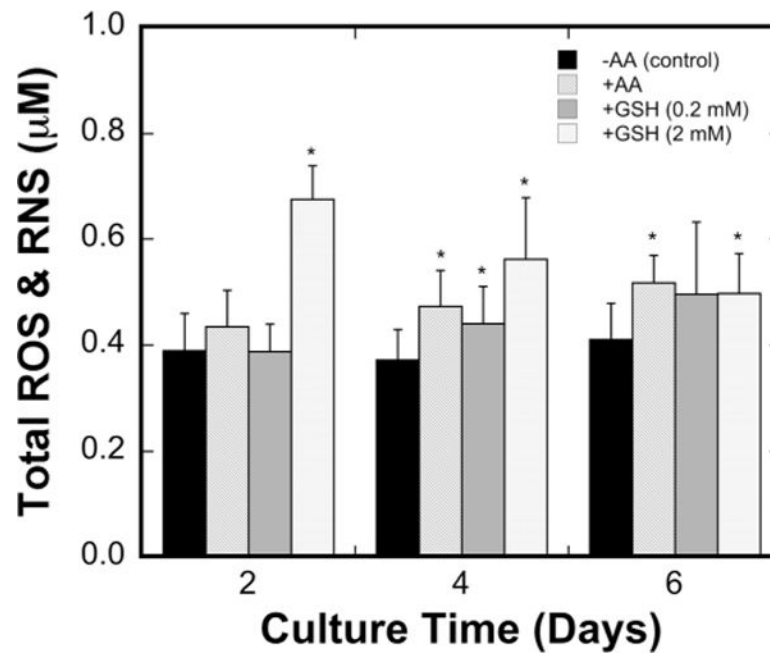


**Figure 2.** Ascorbic acid enhances the survival of primary follicles (60–70 μm). The survival and growth (diameter mean ± standard deviation) of (A and B) 70 μm and (C and D) 60 μm primary follicles cultured in the absence (–AA) and presence (+AA) of ascorbic acid. Significant differences between survival curves were indicated by different lower case letters (a and b,  $P < 0.05$ ). No significant differences in growth were observed ( $P > 0.05$ ). Sample sizes are presented in Table I.



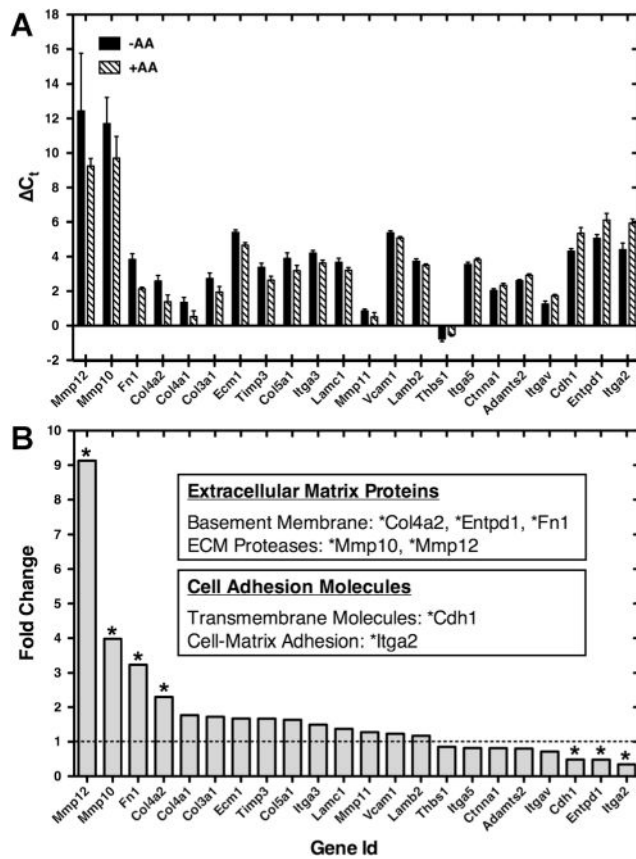
**Figure 3.**

Primary follicle survival in the presence of ascorbic acid (+AA). Representative images of primary follicles (60–70  $\mu\text{m}$ ) cultured in the presence of ascorbic acid on (A) day 0, (B) day 6, (C) day 10, and (D) day 16. Confocal images of (E) growing follicle cultured with ascorbic acid on day 6 and (F) in vivo grown secondary follicle control (red, laminin; green, f-actin, and blue, nucleus). Ascorbic acid significantly increased the survival rate. In the presence of ascorbic acid, primary follicles grew to approximately 250  $\mu\text{m}$  in diameter and developed antral cavities (asterisks). The scale bars represent (A–D) 100  $\mu\text{m}$  and (E) 10  $\mu\text{m}$ .

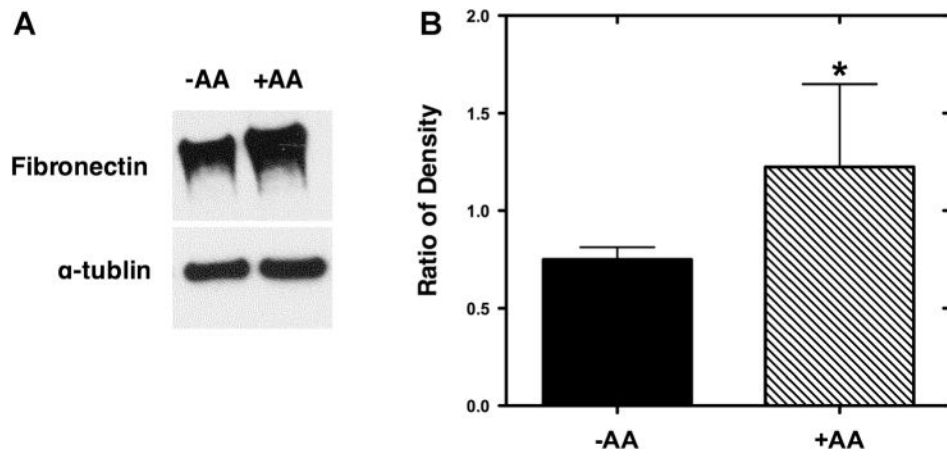


**Figure 4.** Oxidative stress in conditioned medium. Reactive oxygen species (ROS) and reactive nitrogen species (RNS; mean  $\pm$  standard deviation) were measured in the control medium ( $-AA$ ) and medium supplemented with ascorbic acid ( $+AA$ ), 0.2 mM glutathione, and 2 mM glutathione on days 2, 4, and 6 of culture ( $n = 10$  for control and  $n = 12$  for AA/GSH conditions). Asterisks (\*) indicate significant difference relative to control ( $P < 0.05$ ).



**Figure 5.**

Ascorbic acid influences the gene expression of extracellular matrix (ECM) and cell adhesion molecules. (A) Significantly different ( $P < 0.05$ )  $C_t$  values (mean  $\pm$  SD) of extracellular matrix (ECM) and cell adhesion molecules genes in the absence ( $-AA$ ) and presence ( $+AA$ ) of ascorbic acid.  $C_t$  is the number of cycles to exceed the analysis threshold (lower  $C_t$  values represent higher mRNA levels).  $\Delta C_t$  is the difference in gene expression between the gene of interest and a panel of housekeeping genes (*Actb*, *B2m*, and *Gusb*). (B) Fold change of genes in the absence ( $-AA$ ) and presence ( $+AA$ ) of ascorbic acid. Genes up-regulated greater than twofold (or down-regulated less than 0.5) were indicated with asterisks (\*). Fold change was calculated as the normalized gene expression ( $2^{-C_t}$ ) of the  $AA+$  sample divided the normalized gene expression ( $2^{-C_t}$ ) of the  $AA-$  sample.



**Figure 6.**

Western blots confirm fibronectin up-regulation. (A) Western blots were performed on follicles cultured in the absence (-AA) and presence (+AA) of ascorbic acid. To normalize differences in follicle numbers between experiments,  $\alpha$ -tubulin was used as a standard. (B) Densitometry ratios (mean  $\pm$  standard deviation) of fibronectin compared to  $\alpha$ -tubulin ( $n = 3$  for each condition). Significant differences between conditions were indicated via asterisks (\* $P < 0.05$ ).

Table 1

Survival and growth of primary follicles (60–70  $\mu\text{m}$ ).

Size	Condition	Day 0 (n)	Day 2 (n, %)	Day 6 (n, %)	Day 10 (n, %)	Day 14 (n, %)	Day 18 (n, %)
70 $\mu\text{m}$	+AA	48	48 (100%)	40 (83%)	34 (71%)	23 (48%)*	n.d.
	-AA	150	142 (95%)	121 (81%)	87 (58%)	35 (23%)	n.d.
60 $\mu\text{m}$	+AA	42	41 (98%)	28 (67%)	12 (29%)	11 (26%)	7 (17%)*
	-AA	24	23 (96%)	6 (25%)	0 (0%)	0 (0%)	0 (0%)

Size	Condition	Day 0 ( $\mu\text{m}$ )	Day 2 ( $\mu\text{m}$ )	Day 6 ( $\mu\text{m}$ )	Day 10 ( $\mu\text{m}$ )	Day 14 ( $\mu\text{m}$ )	Day 18 ( $\mu\text{m}$ )
70 $\mu\text{m}$	+AA	70 $\pm$ 3	84 $\pm$ 7	106 $\pm$ 15	140 $\pm$ 29	236 $\pm$ 43	n.d.
	-AA	71 $\pm$ 3	91 $\pm$ 10	111 $\pm$ 16	138 $\pm$ 27	251 $\pm$ 47	n.d.
60 $\mu\text{m}$	+AA	61 $\pm$ 3	71 $\pm$ 8	92 $\pm$ 15	133 $\pm$ 27	170 $\pm$ 39	253 $\pm$ 39
	-AA	61 $\pm$ 3	70 $\pm$ 6	82 $\pm$ 11	n.s.	n.s.	n.s.

-AA (70  $\mu\text{m}$ ) data referenced from Tagler et al. (2012).

-AA is control medium or  $\alpha$ MEM supplemented with BSA, fetuin, ITS, and FSH.

Significant differences between conditions were indicated by asterisks ( $P < 0.05$ ).

Diameter presented as mean  $\pm$  standard deviation.

n.d. indicates experiment not done.

n.s. indicates no surviving follicles.

Table II

Gene expression of extracellular matrix (ECM) and cell adhesion molecules.

Gene id	Description	Fold change	95% CI	P-value
Significantly different ( $P < 0.05$ ) and fold change greater than 2 or less than 0.5				
* Mmp12	Matrix metalloproteinase 12	9.13	(0.00001, 29.91)	0.0058
* Mmp10	Matrix metalloproteinase 10	3.98	(0.00001, 9.30)	0.0432
* Fn1	Fibronectin 1	3.24	(2.48, 4.01)	0.0000
* Col4a2	Collagen, type IV, alpha 2	2.31	(1.52, 3.09)	0.0073
* Cdh1	Cadherin 1	0.49	(0.37, 0.60)	0.0004
* Entpd1	Ectonucleoside triphosphate diphosphohydrolase 1	0.48	(0.34, 0.63)	0.0029
* Itga2	Integrin alpha 2	0.35	(0.24, 0.45)	0.0025
Significantly different ( $P < 0.05$ )				
Col4a1	Collagen, type IV, alpha 1	1.77	(1.23, 2.30)	0.0146
Col3a1	Collagen, type III, alpha 1	1.72	(1.18, 2.26)	0.0258
Ecm1	Extracellular matrix protein 1	1.67	(1.43, 1.92)	0.0007
Timp3	Tissue inhibitor of metalloproteinase 3	1.67	(1.29, 2.04)	0.0041
Col5a1	Collagen, type V, alpha 1	1.63	(1.14, 2.13)	0.0142
Itga3	Integrin alpha 3	1.49	(1.26, 1.73)	0.0023
Lamc1	Laminin, gamma 1	1.37	(1.11, 1.64)	0.0137
Mmp11	Matrix metalloproteinase 11	1.28	(1.04, 1.51)	0.0414
Vcam1	Vascular cell adhesion molecule 1	1.23	(1.11, 1.35)	0.0063
Lamb2	Laminin, beta 2	1.17	(1.04, 1.31)	0.0327
Thbs1	Thrombospondin 1	0.85	(0.75, 0.96)	0.0400
Itga5	Integrin alpha 5 (fibronectin receptor alpha)	0.82	(0.73, 0.91)	0.0124
Ctnna1	Catenin (cadherin associated protein), alpha 1	0.81	(0.73, 0.90)	0.0095
Adams2	A disintegrin-like and metalloproteinase (reprolysin type) with thrombospondin type 1 motif, 2	0.80	(0.75, 0.86)	0.0008
Itgav	Integrin alpha V	0.72	(0.62, 0.81)	0.0041
Selected genes not significantly different ( $P > 0.05$ )				
Itgal	Integrin alpha L	18.80	(0.00001, 67.56)	0.1597
Itgam	Integrin alpha M	14.84	(0.00001, 42.17)	0.0870
Itga4	Integrin alpha 4	6.97	(0.00001, 25.77)	0.1044
Mmp9	Matrix metalloproteinase 9	5.48	(0.00001, 14.01)	0.0807
Syt1	Synaptotagmin I	4.96	(0.00001, 15.09)	0.1541
Col2a1	Collagen, type II, alpha 1	3.85	(0.00001, 13.90)	0.1954
Thbs2	Thrombospondin 2	3.28	(0.06, 6.50)	0.0513
Col1a1	Collagen, type I, alpha 1	3.18	(0.76, 5.61)	0.0529
Timp1	Tissue inhibitor of metalloproteinase 1	1.05	(0.88, 1.22)	0.4937

CI, confidence interval.

\* Indicates fold change greater than 2 (or less than 0.5) and statistically significant ( $P < 0.05$ ).