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1 ***Hormone-dependent bacterial growth, persistence and biofilm***
2 ***formation – A pilot study investigating human follicular fluid***
3 ***collected during IVF cycles***

4

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26 **Keywords**

27 Biofilm, human follicular fluid, bacteria, hormones, *in vitro* fertilization

28 **Abstract**

29 Human follicular fluid, considered sterile, is aspirated as part of an *in vitro* fertilization (IVF)
30 cycle. However, it is easily contaminated by the trans-vaginal collection route and little
31 information exists in its potential to support the growth of microorganisms. The objectives of
32 this study were to determine whether human follicular fluid can support bacterial growth over
33 time, whether the steroid hormones estradiol and progesterone (present at high levels within
34 follicular fluid) contribute to the *in vitro* growth of bacterial species, and whether species
35 isolated from follicular fluid form biofilms. We found that bacteria in follicular fluid could
36 persist for at least 28 weeks *in vitro* and that the steroid hormones stimulated the growth of
37 some bacterial species, specifically *Lactobacillus* spp., *Bifidobacterium* spp. *Streptococcus*
38 spp. and *E. coli*. Several species, *Lactobacillus* spp., *Propionibacterium* spp., and
39 *Streptococcus* spp., formed biofilms when incubated in native follicular fluids *in vitro* (18/24,
40 75%). We conclude that bacteria aspirated along with follicular fluid during IVF cycles
41 demonstrate a persistent pattern of growth. This discovery is important since it can offer a
42 new avenue for investigation in infertile couples.

43 **Introduction**

44 Follicular fluid, which surrounds the oocyte during *in vivo* folliculogenesis, is a
45 hypocoagulable, semi-viscous fluid comprising proteins, inorganic compounds,
46 carbohydrates, mucopolysaccharides, lipids, gonadotrophins, steroid hormones,
47 immunoglobulins, cytokines, complement components and growth factors [1,2]. Previous
48 studies have shown that whilst human follicular fluid has antimicrobial properties, it is still
49 capable of supporting microbial growth *in vitro* [3,4]. Cottell *et al.* [5] first reported the
50 presence of bacteria within follicular fluid that had been collected at the time of trans-vaginal
51 oocyte retrieval. In addition, in our recent study, we isolated numerous microorganisms from
52 follicular fluid. The microorganisms were present as asymptomatic colonizers or as
53 contaminants that were introduced into the follicular fluid at the time of trans-vaginal oocyte
54 retrieval [6]. The collected follicular fluids did not appear cloudy/turbid, despite the presence
55 of a mixed microflora, in contrast to acute microbial infections of other body fluids such as
56 urine. In their native environment, the majority of bacteria exist as complex surface-attached
57 communities [7]. This observation prompted our current study to investigate follicular fluid
58 as a medium to support the growth of microorganisms. Since the follicular fluids collected
59 from women undergoing IVF cycles contain high levels of estradiol and progesterone as a
60 result of the IVF stimulation process, we hypothesised that these steroid hormones may affect
61 the growth of microorganisms. We also hypothesised that microorganisms present in human
62 follicular fluids could persist over time and form biofilms in the ovarian follicular fluid.

63

64 **Methods**

65 **1.1 Participants**

66 From September 2007 to November 2008, couples commencing fully stimulated IVF cycles
67 at Wesley-Monash IVF in Brisbane, Australia were invited to enrol in this study. Thirty-six

68 follicular fluid specimens were randomly selected for testing and included equal numbers of
69 clear and blood-stained fluids.

70

71 **1.2 Ethics statement**

72 Ethical approval was obtained from the review boards of Uniting Care Health, Human
73 Research Ethics Committee and Queensland University of Technology Human Ethics
74 Committee. All patients provided informed written consent for their follicular fluids to be
75 used in this study.

76

77 **1.3 Trans-vaginal oocyte retrieval**

78 Follicular fluid was collected by the IVF clinicians at the time of oocyte retrieval as
79 previously described [6]. The IVF unit used a ‘boost’ protocol for controlled ovarian hyper-
80 stimulation. For each study participant, the follicular fluid from the largest most accessible
81 follicle in either the left or the right ovary was aspirated first. Follicular fluid was aspirated
82 directly into sterile test tubes in the operating theatre. The follicular fluid specimens were
83 aseptically transferred to a sterile culture dish to determine if there was an oocyte present.
84 Following removal of the oocyte, the IVF scientists transferred the remaining follicular fluid
85 to a sterile 15 mL Falcon tube for storage at - 80° C. There was significant variability in the
86 volume of follicular fluid collected from each follicle (range < 1 mL – ~12 mL). Therefore,
87 only the follicular fluid samples with a volume of greater than 5 mL were able to be tested
88 using all assays.

89

90 **1.4 Colony identification and 16S ribosomal RNA (rRNA) PCR and sequencing**

91 Isolation and identification of microbial species from the 36 human follicular fluid specimens
92 were performed as previously described [6]. Briefly, calibrated 1 µL inoculating loops were

93 used to inoculate a range of microbiological culture media, which were incubated under 5%
94 CO₂ or anaerobically at 37° C for the isolation and identification of microorganisms.
95 DNA extraction was performed on 1 mL aliquots of each follicular fluid specimen, and then
96 the extracted DNA used as a template for 16S rRNA PCR. The preparation of PCR products
97 for sequencing was performed as per the Australian Genome Research Facility instructions
98 for the preparation of purified DNA (AGRF, St. Lucia, QLD). AGRF sequenced each
99 purified PCR product using a Big Dye 3 sequencing technology (BDT) labelling sequencing
100 platform. The sequence, obtained in a FASTA format was entered into the Basic Local
101 Alignment Search Tool (BLAST, NCBI) for identification of clinical isolates.

102

103 **1.5 Hormonal effect on bacterial growth in follicular fluid**

104 Twelve frozen follicular fluid aliquots were thawed, and 1 µl of each cultured on a range of
105 solid agar plates and in thioglycollate broth containing estradiol and progesterone (Sigma
106 Aldrich, Castle Hill, NSW), at concentrations of 375 µg/L and 800 µg/L respectively (the
107 median concentrations reported in the follicular fluid collected from hyperstimulated women
108 undergoing trans-vaginal oocyte retrieval) [8,9]. Hormones were added to the culture media
109 because these would be present in the native follicular fluids and would degrade over time in
110 culture. Hormones were used only in combination, rather than individually, as both are
111 present at high concentrations in the follicular fluids of women undergoing IVF treatment and
112 trans-vaginal oocyte retrieval. Follicular fluid specimens were also cultured on the same
113 media without the addition of hormones as controls. Agar plates were incubated either
114 aerobically or anaerobically at 37° C. After seven days of incubation at 37° C, positive
115 thioglycollate broths were vortexed and a sterile 1 µL calibrated loop used to subculture broth
116 onto horse blood agar (Oxoid, Adelaide, SA) for quantification and identification of the
117 bacteria present [6].

118 **1.6 Long-term follicular fluid culture**

119 From 24 follicular fluid specimens 1 mL of follicular fluid was aliquoted aseptically into 1.7
120 mL microcentrifuge tubes and incubated at 37° C aerobically. These 24 specimens were
121 vortexed and subcultured daily: 1 µL of the follicular fluid was subcultured, onto horse blood
122 agar and anaerobic blood agar plates (Oxoid) and incubated aerobically and anaerobically at
123 37° C. After (24 hours) incubation, the number of colonies on the horse blood agar were
124 counted and expressed as the number of colony forming units (CFU)/mL of follicular fluid.

125

126 **1.7 Biofilm assay**

127 From the same 24 follicular fluid specimens 300 µL was added to a sterile 13 mm coverslip
128 in a well of a 24-well microtiter plate for the biofilm assay. Triplicate microtiter plates
129 (technical replicates) were prepared and subsequently incubated in aerobic and anaerobic
130 conditions at 37° C. Two coverslips were stained as described below, and the bacteria from
131 the third coverslip were inoculated onto media as described in Methods 1.9. The follicular
132 fluid specimens were incubated within the microtiter plates and left undisturbed for ten days,
133 to mimic the average period of development of follicular fluid within a maturing ovarian
134 follicle [10]. Upon removal, coverslips were gently rinsed with PBS to remove any unbound
135 cells. The coverslips were subsequently placed onto microscope slides and processed using a
136 method adopted from Allison and Sutherland [11]. Briefly, the coverslips were covered with
137 10 mM cetyl pyridinium chloride and air-dried before heat fixation. The biofilms then were
138 stained for 15 minutes with a 2:1 mixture of saturated Congo red (Sigma Aldrich) solution
139 and 10% Tween 20 (Sigma Aldrich). Slides were then rinsed, counterstained with 10% Ziehl
140 carbol fuchsin, rinsed again and dried at 37° C. The prepared coverslips were viewed by light
141 and confocal microscopy.

142

143 Biofilms were viewed by light microscopy using an Olympus BX41 light microscope
144 (Olympus, Tokyo, Japan), and images captured with a MicroPublisher 3.3 RTV camera
145 (Adept Electronic Service, Warriewood, NSW) and QCapture Pro software (QImaging,
146 Surrey, BC). Biofilms were examined under total magnifications of $\times 100$ and $\times 400$.

147

148 Images of the biofilms also were acquired by a TCS SP5 confocal laser scanning microscope
149 (Leica Microsystems, Germany) equipped with a Leica HCX PL APO CS $\times 10$ objective, and
150 a Leica HCX PL APO CS $\times 63$ oil immersion objective (NA 1.4). To visualise the Congo
151 red and carbol fuschin signal, the excitation wavelength was set at 561 nm, and the
152 fluorescence emission was detected between 567- 668 nm. A series of z stack images were
153 acquired through a volume of 60 μm and the images analysed using LAS AF (Leica
154 Microsystems).

155

156 Biofilm maturity was graded based on the presence of key characteristic and structural
157 features described previously by Simmons *et al.* [12]. Biofilms were classified as grade I—
158 planktonic cells (isolated free floating cells, not adherent to the slide) and cells adherent to
159 the conditioning film; grade II—microcolonies and groups of cells (most likely planktonic)
160 attached to each other; grade III—extending/growing microcolonies (towers) with
161 extracellular matrix fibrils, creating interconnections between the microcolonies, thus giving
162 a cobweb appearance; and grade IV—towers with subterranean channels and amorphous
163 extracellular material, giving a honeycomb appearance between the microcolonies.

164

165 **1.8 Biofilm assay – scanning electron microscopy (SEM)**

166 SEM was used to visualise biofilm production for four of the follicular fluid specimens and
167 two ATCC controls (*Lactobacillus gasseri* and *Bacteroides fragilis*). 300 μL of each of the

168 follicular fluid specimens or ATCC overnight broth cultures was added to a single sterile
169 13mm coverslip in each well of a 24-well microtiter plate. Plates were performed in
170 duplicate so that one set was grown for 5 days, and the other 10 days. Glass coverslips were
171 fixed in 3% glutaraldehyde in 0.1 M sodium cacodylate buffer, pH 7.4 for 24 hours at 4° C.
172 After primary fixation, the coverslips were washed three times for 10 minutes in 0.1 M
173 cacodylate buffer and post fixed for one hour in either 1% Osmium tetroxide or 1% Osmium
174 tetroxide with 0.15% ruthenium red. Post-fixation, coverslips were twice washed in water and
175 dehydrated in an ascending ethanol series (50, 70, 90, and 100% (twice)) before drying with
176 100% hexamethyldisilazane (HMDS) (Sigma Aldrich). Specimens were mounted on
177 aluminium stubs with adhesive carbon tape and then sputter coated with 10 nm of gold
178 (Leica SCD005 sputter coater). Examination of samples was performed using a FEI Quanta
179 3D Focused Ion Beam SEM, operating at 10 kv.

180

181 **1.9 Biofilm culture**

182 The third coverslip (from Methods 1.7) was rinsed with sterile PBS; a sterile swab was used
183 to remove the biofilm from the coverslip surface. This bacterial suspension was made in
184 thioglycollate broth; a 1 µL calibrated inoculating loop was used to subculture the biofilm
185 onto horse blood agar and anaerobic blood agar plates (Oxoid). Plates were incubated
186 aerobically at 37° C for 24-48 hours and at 37° C for up to 7 days. Isolated bacterial species
187 were identified as previously described [6].

188

189 **Results**

190 **Follicular fluid culture and bacterial colony identification**

191 Bacteria were cultured from each of the 24 follicular fluid specimens tested. A single
192 bacterial species was isolated from 15/24 (63%) of the specimens, of which 7/15 (47%)

193 contained *Lactobacillus* species. *Propionibacterium* spp. (n = 5), *Peptostreptococcus* spp. (n
194 = 2), or *Salmonella enterica* (n = 1) were isolated in the remaining specimens. Two bacterial
195 species were isolated from 6/24 (25%) specimens, and five of these (83%) contained
196 *Lactobacillus* spp. Only 3/24 (12%) follicular fluid specimens contained three bacterial
197 species and *Lactobacillus* spp. was isolated from each of these specimens. *Lactobacillus* spp.
198 and *Propionibacterium* spp. were the most prevalent isolates detected in follicular fluid,
199 isolated from 51% and 14% specimens respectively (Table 1). Only 13/24 follicular fluid
200 samples were of a sufficient volume for the complete analyses.

201

202 **Hormonal modulators of bacterial growth in solid media and broths**

203 When follicular fluid was cultured on solid agar media and incubated under appropriate
204 atmospheric conditions, with or without hormone supplements, there were no differences in
205 the number of isolated CFUs/mL (10^3 - $> 10^6$). However, differences were observed in the
206 growth patterns of bacteria cultured in thioglycollate broth (with and without hormones) for
207 5/12 follicular fluid specimens tested (Table 2). *Lactobacillus* spp. were recovered from 7/12
208 (58%) thioglycollate broths. For one specimen (see Table 2, No. 12), *Lactobacillus crispatus*
209 and *L. gasseri* were cultured from the hormone-supplemented thioglycollate media, but only
210 *L. crispatus* was recovered from the hormone-free media. However, for the remaining
211 *Lactobacillus* containing specimens, there were slight differences in the CFUs/mL of the
212 *Lactobacillus* spp. isolated in the presence or absence of hormones. *Bifidobacterium* spp. was
213 detected in two follicular fluid specimens, and where the supplemental hormones supported
214 the growth of the *Bifidobacterium* spp., this bacterium did not grow in the absence of
215 hormones. In contrast, the supplemental hormones inhibited the growth of *E. coli* and *S.*
216 *agalactiae*, but growth was observed within the thioglycollate broth, without the addition of
217 exogenous steroid hormones (see Table 2).

218 **Long-term follicular fluid *in vitro* culture**

219 All follicular fluid specimens cultured *in vitro* demonstrated increasing numbers of CFUs/mL
220 at each subculture until eight days post-incubation ($< 10^3 - > 10^6$), after which the number of
221 cultivable bacteria reached a plateau and the numbers for these bacteria remained constant for
222 the remaining 27 weeks ($10^3 - 10^6$). Furthermore, after 5-8 days incubation, only a single
223 bacterial species in pure culture could be isolated from all follicular fluid specimens. With the
224 exception of *S. enterica*, all of the bacterial species isolated after eight days were Gram-
225 positive (*Lactobacillus* spp., *Bifidobacterium* spp. and *Staphylococcus* spp.). Viable Gram-
226 positive bacteria could be recovered from these follicular fluids, which were incubated *in*
227 *vitro* for 28 weeks.

228

229 **Biofilm assay**

230 The 24 follicular fluid specimens were tested to determine their ability to form biofilms on
231 glass coverslips in wells of a 24-well plate. Biofilms formed *in vitro* after incubation at 37° C
232 for 18/24 (75%) of follicular fluids. Of these biofilms, 14 were monomicrobial and four were
233 polymicrobial as determined by the number of different colony types identified by traditional
234 microbiological culture. The bacterial species *L. gasseri*, *L. crispatus*, *Bifidobacterium*
235 *longum*, *S. agalactiae*, *S. anginosus* and *S. entericus*, if initially present in follicular fluid,
236 were always recovered from the biofilm well sub-cultures after 10 days of *in vitro* culture.
237 Other bacterial species, including CoNS, *Peptostreptococcus* spp. and *E. coli*, were only
238 cultured from the original follicular fluid specimens. In 16/24 (66%) of the follicular fluids
239 tested, the primary culture demonstrated polymicrobial colonisation, but only a single species
240 was detected after 10 days of incubation within each well of the microtiter plate. In addition,
241 if only a single species was detected in the primary follicular fluid culture, the same single
242 species was always detected in the microtiter well culture. The facultative anaerobic species,

243 *Staphylococcus* spp., *Streptococcus* spp. and *Lactobacillus* spp., were isolated following
244 incubation under both aerobic and anaerobic conditions. In contrast, the strict anaerobes
245 *Bifidobacterium* spp., and *Peptostreptococcus* spp. were isolated only when the follicular
246 fluid specimens were incubated anaerobically.

247

248 For six biofilms slides, there was no evidence of biofilm formation or planktonic cell
249 adhesion. Instead, the Congo red stain revealed the presence of a polysaccharide-containing
250 conditioning film. The images shown in Figure 1 are representative of the different types of
251 *Lactobacillus* spp. biofilms observed for follicular fluids incubated both under aerobic and
252 anaerobic atmospheric conditions. Variations in the architecture of mature biofilms was
253 observed, ranging from flat homogenous cell layers through to cell clusters, microcolonies
254 and towers with channelling and significant quantities of amorphous extracellular material
255 between the more complex heterogeneous structures (Figure 1 B-D). The simplest biofilm
256 architecture visualised by light microscopy was seen for follicular fluid cultures that were
257 incubated aerobically. The grade II biofilm presented in Figure 1A (b) demonstrated few
258 microcolonies (which proliferate at fixed positions [13,14], and some of these extended to
259 form towers (outward growing masses of bacterial cells). Initial microcolonies were formed
260 by cells growing outwards into towers, with the highest points of elevation appearing white in
261 greyscale using confocal microscopy, as shown in Figure 2A (a) and (b). A grade III biofilm
262 was characterised by fibril formation, as observed by light microscopy, following anaerobic
263 incubation (Figure 1B (b)); with a carpet-pile appearance as observed with confocal
264 microscopy with some microcolonies extending to form towers, as shown in Figure 2B (a).
265 The polymeric extracellular matrix appeared to be more abundant in microcolonies (Figure
266 1C (b)). When magnified ($\times 4$ zoom, confocal microscopy), crater-like formations were
267 visible (Figure 2B (b)). In the grade IV biofilm grown under anaerobic conditions, complex

268 networks were observed between the microcolonies and towers and these had a cobwebbed
269 appearance, as shown in Figure 2C (a) and (b). Grade IV biofilms were also visualised using
270 light microscopy (Figure 1D (b)) and an amorphous polymeric extracellular matrix was
271 observed surrounding the interconnecting microcolonies, towers and cells. In these biofilms,
272 continuous layers of cells covered the coverslip, with clusters forming prominences. These
273 biofilms had a honeycomb appearance, as shown in Figure 2D (a) and (b) and could be
274 visualised using three dimensional and orthogonal confocal images. This image (Figure 3)
275 demonstrated the presence of channels within the biofilms and hollow areas under the towers.
276 All observed biofilms showed an uneven spatial distribution, which is consistent with the
277 previously described characteristic of *in vitro* biofilms [15].

278

279 **Biofilm assay – scanning electron microscopy**

280 Scanning electron microscopy (SEM) was used to visualize the surface topography of
281 biofilms grown *in vitro* on glass coverslips for 5 days or 10 days. SEM revealed the presence
282 of bacterial cell aggregates covered by a thin layer of glycocalyx (Figure 4A (i)), as well as
283 microcolonies (Figure 4A (ii)). The surface topography, with a crater-like appearance was
284 similar to that seen in images obtained using confocal microscopy (Figure 2B). The
285 preservation of the glycocalyx was enhanced when the post-fixative OsO₄ (Figure 4C (i)) was
286 combined with ruthenium red (Figure 4C (ii)). The use of the ruthenium red enhanced the
287 stabilization and visualization of the highly charged anionic mucopolysaccharides forming
288 the extracellular glycocalyx. Evidence of biofilm maturity, extensive extracellular
289 polysaccharide and microcolonies was apparent in both five and ten day-old biofilms.

290

291 **Discussion**

292 Follicular fluids collected at the time of trans-vaginal oocyte retrieval were found to harbour
293 bacteria and continued to support the bacterial growth *in vitro*. These results demonstrate that
294 bacterial species that colonise follicular fluid, or gain access to the follicular fluid at the time
295 of oocyte retrieval, may be a relevant focus of infertility investigations in couples with
296 idiopathic infertility or repeated adverse IVF treatment outcomes.

297

298 Our study demonstrated that *in vitro*, elevated levels of estradiol and progesterone in
299 thioglycollate broth media enhanced the growth of the high numbers of CFUs of
300 *Lactobacillus* spp., which were originally present within follicular fluid specimens. This is
301 consistent with *Lactobacillus* spp. growth in the lower genital tract. At puberty, as estrogen
302 levels increase, the vaginal pH drops and *Lactobacillus* spp. dominate the vaginal microflora
303 [16]. After menopause, the vaginal microflora reverts to the pre-menarchal state and the
304 *Lactobacillus* spp. decline [17]. However, if post-menopausal women receive estrogen
305 replacement therapy, the vaginal pH decreases and the concentration of *Lactobacillus* spp.
306 increases [18]. The effect of endogenous steroid hormones on lactobacilli has previously been
307 monitored in IVF patients. Jakobsson and Forsum [19] reported that during IVF treatment,
308 with increasing estrogen levels, three major vaginal lactobacilli (*L. crispatus*, *L. gasseri* or *L.*
309 *jensenii*) were predominant. These results are consistent with the findings reported in this
310 paper, as these species were the only lactobacilli isolated from follicular fluid (collected prior
311 to ovulation when estradiol levels were highest), although *Bifidobacterium* spp. was also
312 isolated from the hormone-supplemented media. Both genera can metabolise carbohydrates
313 from glycogen degradation in response to elevated estradiol levels, making the female genital
314 tract a niche environment, enhancing the ability of the lactobacilli to persist and protect the
315 genital tract epithelium from opportunistic infection [20]. *Bifidobacterium* spp., whilst

316 traditionally accepted as members of the normal regional flora of the gastrointestinal tract,
317 have more recently been detected in the vaginal flora of healthy women [20,21,22]

318

319 During an IVF treatment cycle, hormones including follicle stimulating hormone and human
320 chorionic gonadotropin are administered to trigger the simultaneous maturation of multiple
321 ovarian follicles. This results in the production of elevated levels of estradiol and
322 progesterone within the follicular fluids collected from women undergoing trans-vaginal
323 oocyte retrieval for IVF at levels approximately eight and three times higher, respectively,
324 than those of women with normal cycles [8,9]. The ovarian granulosa cells are the dominant
325 source of these steroid hormones. Granulosa cells may be aspirated with the follicular fluid
326 and the oocyte at the time of trans-vaginal oocyte retrieval, however once removed from the
327 ovary, the granulosa cells are no longer under pituitary control and so rapidly proceed to
328 undergo apoptosis [23,24], and do not continue to produce the steroid hormones *in vitro* as
329 they do *in vivo* [25]. Thus, over time in culture without the addition of exogenous hormones,
330 the impact of steroid hormone levels on microbial growth would not be observed.

331

332 Studies have demonstrated that the hormone concentration within maturing follicles is
333 significantly higher than that found in the systemic circulation, as it is governed by the
334 hyaluronan composed cumulus-oocyte complex [2]. In the current study, hyaluronidase-
335 producing species (*Propionibacterium* spp., *Streptococcus* spp. and *E. coli*) were isolated
336 from follicular fluid. The hyaluronidase virulence factor of these species could lead to poor
337 quality cumulus cells via enzymatic hyaluronan degradation, decreased steroid hormone
338 synthesis and ultimately to detrimental effects of these more pathogenic bacteria within the
339 follicle. Our findings indicate that high concentrations of steroid hormones within ovarian
340 follicular fluid can influence bacterial growth in a species-dependent manner.

341 We demonstrated that follicular fluid supported the survival of viable bacteria for a period of
342 at least 28 weeks. This growth occurred without the addition of supplemental nutrients or the
343 removal of metabolic waste products; such processes that would occur *in vivo* and may
344 facilitate continual microbial colonisation of the follicular fluid. Gurgan *et al.* [4] found that
345 filter-sterilised and centrifuged follicular fluid was inhibitory to the Gram-positive bacteria *S.*
346 *aureus*, *S. agalactiae* and *L. monocytogenes*, as these species did not survive *in vitro* for
347 longer than four days. However, Gurgan *et al.* [4] did demonstrate that follicular fluid
348 supported the growth of a range of Gram-negative species and the yeast *Candida albicans* for
349 up to 15 days post-inoculation. These results are inconsistent with those reported here, where
350 predominantly Gram-positive species survived, most likely as a result of the composition of
351 the follicular fluid, which may have been significantly altered by the filtration and
352 centrifugation prior to inoculation. To the best of our knowledge, our study is the first
353 continuous follicular fluid culture experiment lasting 28 weeks, indicating that follicular fluid
354 is an excellent growth medium for microorganisms.

355

356 Many of the bacteria that we detected within follicular fluid have previously been shown to
357 form biofilms *in vitro* when cultured on vaginal epithelial cells [26], *in vivo* on the surface of
358 intrauterine devices [27] and in amniotic fluid sludge aspirated from women diagnosed with
359 intrauterine infection [28]. However, despite the range and load of bacteria isolated from
360 follicular fluid, this fluid did not have a turbid appearance, which may suggest that bacteria
361 within the follicle were present predominantly in biofilms, rather than as planktonic bacteria.
362 Thus, the process of trans-vaginal oocyte retrieval may result in puncture and collection of
363 the follicle wall biofilm within the lumen of the aspiration needle, and the subsequent transfer
364 of biofilm components, including bacteria into the follicular fluid at the time of egg retrieval.

365

366 The majority of mature biofilms in the present study revealed only a single cultivable species
367 after 10 days, even when two different species of lactobacilli were initially isolated from the
368 follicular fluid prior to *in vitro* incubation. This may be due to competition amongst the
369 lactobacilli themselves, or because of alterations in adhesion. It has been reported that some
370 bacterial species can co-colonise the surface of urogenital tract cells already colonised by
371 lactobacilli; however, after lactobacilli are established, they can cause other species of
372 bacterial cells to detach from the epithelium, which may offer a defence mechanism by
373 preventing attachment of pathogens to the epithelium [29,30]. In polymicrobial biofilms,
374 early colonising species often promote the establishment of other species [31,32]; however,
375 whilst the initial interactions may be synergistic, once the biofilm is established, competition
376 between species can result in the dominance of a single species [33]. Quorum sensing, the
377 process by which microorganisms communicate within a population, establishes the overall
378 population size of each representative microbial species and initiates alterations in gene
379 expression . In addition to quorum sensing, microbial growth within the biofilm is also
380 modulated by these metabolic cues [34]. Species production of lactic acid, succinic acid and
381 isobutyric acid has been reported to enhance both the synergistic and competitive interactions
382 between different microbial species within a biofilm. Acid production resulting in a pH shift,
383 often causes a reduction in the microbial diversity, as many species are acid intolerant [35].
384 We hypothesise that the predominance of monomicrobial *in vitro* biofilms reported in this
385 study might therefore be due to the lactic acid and H₂O₂ production by *Lactobacillus* spp.
386
387 Studies of biofilms have reported that bacteria can exist in a viable, but non-cultivable state
388 [36]. It is therefore possible that cultures of 10-day biofilms detected only those species that
389 were actively replicating, whereas molecular techniques targeting 16 S rRNA would be able
390 to detect and quantify all species (both viable and non-viable) present within the biofilm. This

391 study did report that one culture-negative follicular fluid sample did not produce any visible
392 growth on the range of solid agar or in the thioglycollate broth; however sequencing of the 16
393 S rRNA PCR product identified a sequence matching an uncultured bacterial clone, again
394 highlighting the need for molecular based techniques. To gain further understanding of an
395 individual species contribution to a biofilm, quantitative 16 S rRNA PCR assays could be
396 used to test cultures at various time points, in order to establish growth curves for each
397 individual species and compare culture results to PCR assay results to identify early and late
398 species contributing to the biofilm.

399

400 An ovarian follicle has all the characteristics necessary to support the development of a
401 microbial biofilm. Bacterial cells that would contribute to the formation of the biofilm have
402 been detected in this work (see Table 1 and Figure 1). On some biofilm slides there is
403 evidence of polysaccharide accumulation (in the absence of bacterial cells), which shows that
404 there are components of follicular fluid (substrate) that could form a conditioning film to
405 allow microbial attachment to the inside wall of the follicle (substratum). Furthermore, the *in*
406 *vitro* development of various grades of biofilms in human follicular fluid (Figure 1) is
407 demonstrated. Mature biofilms are characterised by the development of towers and fluid-
408 filled channels [37,38]. Mature biofilms reportedly exhibit a variety of phenotypes due to the
409 changes in their three-dimensional structure, occurring in response to changes in cell density,
410 osmolarity, temperature, pH and nutrient supply [36]. We showed that even follicular fluids
411 containing the same bacterial species demonstrated different grades of biofilms and different
412 morphology.

413

414 On six biofilm slides, no bacteria were observed, only a conditioning film was present. This
415 could be due to the removal of any bacterial cells (by the PBS rinse prior to staining) present

416 in the early stage of biofilm formation prior to permanent attachment to the conditioning film.
417 Alternatively, the follicle may have been colonised by non-biofilm forming bacterial species.
418 Further studies examining the effect of these biofilm bacterial species on each other would
419 therefore, be beneficial to our understanding of the impact of single species and
420 polymicrobial biofilms on reproductive health outcomes.

421

422 The findings reported in this paper have shown that (1) the ovarian steroid hormones
423 (estradiol and progesterone) are capable of modulating the *in vitro* growth of some microbial
424 species; (2) follicular fluid supports the long-term survival of microorganisms; and (3)
425 bacteria isolated from follicular fluid can form biofilms *in vitro*. Further characterisation of
426 the microorganisms detected in follicular fluid and their metabolites will increase our
427 understanding of the effects of follicular fluid microorganisms on oocyte quality, on IVF
428 fertilisation and pregnancy rates, and on early pregnancy events. Knowledge of the
429 microorganisms present within this anatomical niche may lead to an improved understanding
430 of ovarian resistance to infection and also on IVF outcomes.

431

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467 **Table 1 Bacterial genera isolated and identified from cultures of follicular fluid**

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Genus and species	Follicular fluid culture n = 24	Percentage of total n = 35 isolates ³	Biofilm culture n = 18	Percentage of total n = 23 isolates ³
<i>Lactobacillus gasseri</i> ¹	9	27%	9	40%
<i>L. crispatus</i> ¹	7	20%	7	30%
<i>L. jensenii</i> ¹	2	5%	1	5%
CoNS ²	3	9%	0	0%
<i>Propionibacterium</i> spp.	5	14%		
<i>Peptostreptococcus</i> spp.	2	5%	0	0%
<i>B. longum</i>	2	5%	1	5%
<i>S. agalactiae</i>	1	3%	1	5%
<i>S. anginosus</i>	1	3%	1	5%
<i>Micrococcus</i> spp.	1	3%	1	5%
<i>Salmonella enterica</i>	1	3%	1	5%
<i>Escherichia coli</i>	1	3%	0	0%
Total number of isolates	35		23	

469 ¹*Lactobacillus* spp. were the most prevalent bacteria in follicular fluid 18/35 (51%); ²CoNS coagulase negative470 staphylococci; ³ Some follicular fluids contained more than one bacterial species, giving the total number of

471 isolates as greater than the number of follicular fluids tested.

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Table 2 Bacterial growth in hormone supplemented thioglycollate broth

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Specimen	Species identified	CFU/mL Hormone supplemented media ²	CFU/mL Hormone-free media ²
1	<i>Bifidobacterium</i> spp.	10 ⁴	No growth
2	<i>Bifidobacterium</i> spp.	10 ⁴	No growth
3	<i>E. coli</i>	No growth	<10 ³
4	<i>L. crispatus</i>	10 ⁶	10 ⁴
5	<i>L. crispatus</i> , <i>L. gasseri</i>	<10 ^{3*}	<10 ³
6	<i>L. crispatus</i> , <i>L. gasseri</i>	10 ⁴	<i>L. crispatus</i> only <10 ³
7	<i>L. gasseri</i>	10 ⁶	10 ⁶
8	<i>L. gasseri</i>	10 ⁶	10 ⁴
9	<i>L. gasseri</i>	10 ⁶	10 ⁶
10	<i>L. jensenii</i>	10 ⁴	10 ⁴
11	No growth ¹		
12	<i>S. agalactiae</i>	No growth	<10 ³

¹ Culture negative, however positive by 16 S rRNA PCR assay, ² Thioglycollate broths, * <10³ represents growth in thioglycollate broth but not on solid agar subculture plates

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485 **Figure legends**

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487 Figure 1. Light micrographs of Gram stains and biofilm slides.

488 A – D (a)) Light microscopy image at $\times 1000$ total magnification of Gram stained bacterial *S.*

489 *agalactiae* and *Lactobacillus* spp. colonies cultured from biofilms. (A - D (b)) Light

490 microscopy image at $\times 1000$ total magnification of Congo red stained *S. agalactiae* and

491 *Lactobacillus* spp. biofilms grown for 10 days on glass coverslips.

492

493 Figure 2. Laser scanning confocal microscopy images of biofilms.

494 (A – D (a)) Laser scanning confocal microscopy image of various grades of biofilms at \times

495 630 total magnification. Scale bars represent 25 μM (A – D (a) Laser scanning confocal

496 microscopy images of various grades of biofilms at $\times 630$ total magnifications plus $\times 4$ zoom.

497 Scale bars represent 7.5 μM (C (b)) Image of cobwebbing. (D (b)) Image of the

498 honeycombed region. The arrow points to the cavities in the honeycombs. Grade II biofilms

499 presented in Figure 2 A (a and b) demonstrated few microcolonies, some of which extended

500 to form towers. A grade III biofilm was characterised by a carpet-pile appearance with some

501 microcolonies extending to form towers (Figure 2B (a)). Grade IV biofilms had a honeycomb

502 appearance where the continuous layers of cells covered the coverslip with clusters forming

503 prominences (Figure 2D (a) and (b)).

504

505 Figure 3. Representative orthogonal view of a 10-day-old biofilm incubated under anaerobic

506 conditions.

507 Laser scanning confocal microscopy image of a mature Grade IV biofilm at $\times 630$ total

508 magnifications plus $\times 4$ zoom. The orthogonal view allows the representation of the 3D

509 biofilm to be presented in 2D. The cross hairs indicate the area of the biofilm presented in 2D

510 along the borders of the image. Depicted below and to the right of the main image are the yz
511 and xz planes respectively. In this biofilm, there are channels between the microcolonies
512 appearing in ‘gaps’ in the 2D structure and indicated by the white arrows. The hollow
513 interconnecting areas under the towers are indicated by the black arrow.

514

515 Figure 4. Scanning electron micrographs of 10-day-old biofilms.

516 Individual bacilli, identified as *Lactobacillus* spp. (day 5) (arrows) are covered by a light
517 glycocalyx (4A (i)) and a control ATCC strain *Bacteroides* spp. (day 5) by a more mature
518 biofilm (4A (ii)) both captured at $\times 10000$ magnification. Figures 4B (i (day 5) and ii (day
519 10)) represent osmium fixed biofilms containing coccoid bacteria identified as *Streptococcus*
520 spp. (by culture) (arrows) at $\times 15000$ and $\times 30000$ magnification. Glycocalyx is not well
521 preserved using traditional aldehyde fixatives followed by OsO_4 post-fixation. Figure 4C (i)
522 (day 10) was fixed using OsO_4 , which stabilises lipids and figure 4C (ii) (day 10) was fixed
523 using an aldehyde fixative followed by OsO_4 plus ruthenium red to enhance the preservation
524 of the anionic polysaccharides in the glycocalyx.

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