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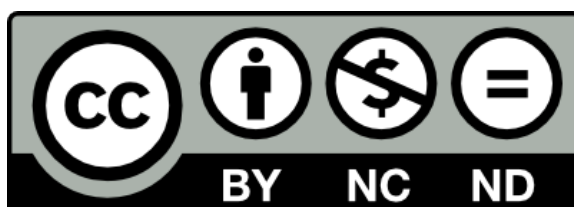
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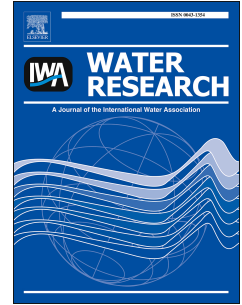


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It's official – *Cryptosporidium* is a gregarine: What are the implications for the water industry?

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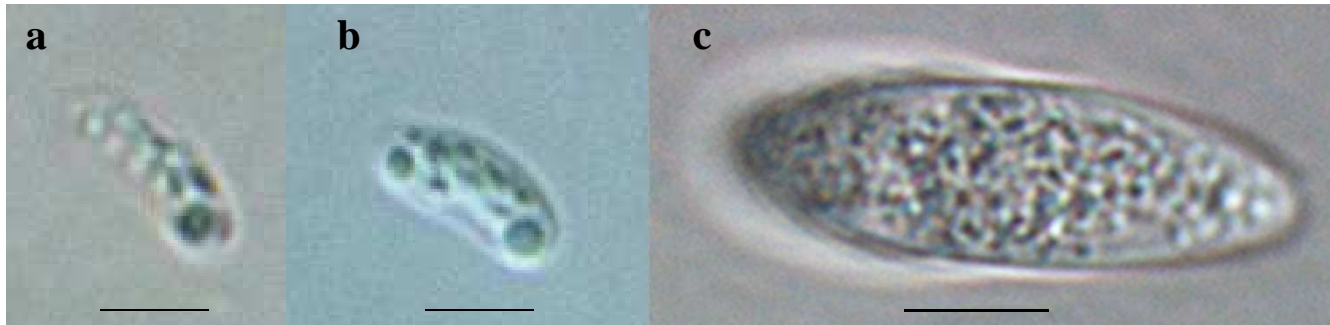
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Novel *Cryptosporidium* life cycle stages



1 Review

2

3 **It's official – *Cryptosporidium* is a gregarine: what are the implications for the water**
4 **industry?**

5

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18

19

20 **Abstract**

21

22 Parasites of the genus *Cryptosporidium* are a major cause of diarrhoea and ill-health in
23 humans and animals and are frequent causes of waterborne outbreaks. Until recently, it was thought
24 that *Cryptosporidium* was an obligate intracellular parasite that only replicated within a suitable
25 host, and that faecally shed oocysts could survive in the environment but could not multiply. In
26 light of extensive biological and molecular data, including the ability of *Cryptosporidium* to
27 complete its life cycle in the absence of a host and the production of novel extracellular stages,
28 *Cryptosporidium* has been formally transferred from the Coccidia, to a new subclass,
29 Cryptogregaria, with gregarine parasites. In this review, we discuss the close relationship between
30 *Cryptosporidium* and gregarines and discuss the implications for the water industry.

31

32 **Keywords:** *Cryptosporidium*; gregarine; cell-free; gamont-like extracellular stages; water industry;

33

34 1 Introduction

35

36 The Apicomplexan parasite *Cryptosporidium* is a major cause of severe diarrhoea,
37 developmental problems and death in young children and chronic, life-threatening disease in
38 immunocompromised and malnourished individuals (Guerrant et al., 1999; Snelling et al., 2007;
39 Costa et al., 2011; Kotloff et al., 2013; Striepen, 2013). No vaccines are available for
40 *Cryptosporidium* (Mead, 2014) and current treatment options for cryptosporidiosis are limited, with
41 only one drug, nitazoxanide (NTZ), exhibiting moderate clinical efficacy in children and
42 immunocompetent people, and none in people with HIV (Abubakar et al., 2007; Amadi et al., 2009).
43 Of the 31 valid species (Costa et al., 2016; Li et al., 2015; Ryan et al., 2015; Holubová et al., 2016;
44 Kváč et al., 2016; Zahedi et al., 2016), *Cryptosporidium parvum* and *Cryptosporidium hominis* are
45 responsible for the majority of human infections, although in some countries, *C. meleagridis* is as
46 prevalent as *C. parvum* in human populations (Xiao, 2010).

47 Transmission of the parasite occurs via the faecal-oral route, either by ingestion of
48 contaminated water or food, or by human-to-human or animal-to-human transmission (Xiao, 2010).
49 The World Health Organization has categorized *Cryptosporidium* as a reference pathogen for the
50 assessment of drinking water quality (Medema et al., 2006). This is because oocysts produced by
51 *Cryptosporidium* are extremely hardy, easily spread via water, resistant to inactivation by chlorine
52 and are difficult to remove from drinking water, without the use of expensive and lengthy filtration
53 (Jakubowski, 1995; Striepen and Kissinger, 2004).

54 Waterborne transmission is a major mode of transmission and *Cryptosporidium* was the
55 etiological agent in 60.3% (120) of the waterborne protozoan parasitic outbreaks that have been
56 reported worldwide between 2004 and 2010 (Baldursson and Karanis, 2011). The severity of
57 infections vary, depending on the species involved, but for zoonotic species, the dose required to
58 cause an infection in 50% of subjects (ID₅₀) is estimated to be 10–83 oocysts for *C. hominis* and

59 132 for *C. parvum* (DuPont et al., 1995; Okhuysen et al., 1998; Chappell et al., 2006). The
60 minimum infectious dose for *C. meleagridis* has yet to be determined (Chappell et al., 2011).
61 Although the lowest infectious dose for *C. hominis* has been calculated to be 10 oocysts, in reality,
62 one oocyst could be sufficient to cause infection in humans through direct or indirect routes of
63 transmission (Chappell et al., 2006).

64 In addition to the apical complex, one main and unique feature of the phylum Apicomplexa,
65 to which *Cryptosporidium* belongs, is the widespread presence of the apicoplast. This four-
66 membrane-encased relict plastid (35kb genome) of secondary endosymbiotic origin is thought to
67 have originated by engulfment of a chloroplast-containing alga by the primitive eukaryotic ancestor
68 of the Apicomplexa (Lim and McFadden, 2010). Microscopic, molecular, genomic and biochemical
69 data indicate that *Cryptosporidium* differs from other apicomplexans in that it has lost the
70 apicoplast (like the colpodellids and other gregarines) (rev. in Lim and McFadden, 2010), as well as
71 the genomes for both the plastid and the mitochondrion (Zhu et al., 2000; Abrahamsen et al., 2004;
72 Xu et al., 2004). *Cryptosporidium* also differs from other apicomplexans in fundamental features
73 such as motility and invasion (Wetzel et al., 2005).

74 Until recently, *Cryptosporidium* was classified as a coccidian parasite. However, it has long
75 been speculated that *Cryptosporidium* represents a ‘missing link’ between the more primitive
76 gregarine parasites and coccidians. The similarities between *Cryptosporidium* and gregarines have
77 been supported by extensive microscopic, molecular, genomic and biochemical data (Pohlenz et al.,
78 1978; Bull et al., 1998; Carreno et al., 1999; Beyer et al., 2000; Hijjawi et al., 2002; Leander et al.,
79 2003a; Hijjawi et al., 2004; Rosales et al., 2005; Barta and Thompson, 2006; Butaeva et al., 2006;
80 Valigurová et al., 2007; Boxell et al., 2008; Karanis et al., 2008; Zhang et al., 2009; Borowski et al.,
81 2008; 2010; Hijjawi, 2010; Hijjawi et al., 2010; Templeton et al., 2010; Karanis and Aldeyarbi,
82 2011; Boxell, 2012; Koh et al., 2013, 2014; Huang et al., 2014; 2015; Clode et al., 2015;
83 Valigurová et al., 2015; Aldeyarbi and Karanis, 2016a; 2016b; 2016c; Edwinson et al., 2016;

84 Paziewska-Harris et al., 2016), which have served as the basis for the formal transfer of
85 *Cryptosporidium* from subclass Coccidia, class Coccidiomorpha to a new subclass, Cryptogregaria,
86 within class Gregarinomorpha (Cavalier-Smith, 2014). The genus *Cryptosporidium* is currently the
87 sole member of Cryptogregaria and is described as comprising epicellular parasites of vertebrates
88 possessing a gregarine-like feeder organelle but lacking an apicoplast (Cavalier-Smith, 2014).
89 According to the International Code of Zoological Nomenclature (ICZN)
90 (<http://www.iczn.org/iczn/index.jsp>), once a species has been formally re-classified in a peer-
91 reviewed publically available journal, then that re-classification stands (unless challenged in the
92 literature). As this re-classification has not been challenged, *Cryptosporidium* is now officially a
93 gregarine.

94

95 **2.0 What are gregarines?**

96

97 Gregarines (phylum Apicomplexa; class Gregarinomorpha) are a very diverse group of
98 large, single-celled “primitive” apicomplexan parasites that primarily infect the intestines and other
99 extracellular spaces of invertebrates and lower vertebrates (mainly arthropods, molluscs and
100 annelids), which are abundant in natural water sources (Leander et al., 2003a; 2003b; Barta and
101 Thompson, 2006; Leander, 2007; Valigurová et al., 2007). The transmission of gregarines to new
102 hosts usually takes place by oral ingestion of oocysts in both aquatic and terrestrial environments.
103 Four or more sporozoites (depending on the species) escape from the oocysts, find their way to the
104 appropriate body cavity and attach to, or penetrate, the host cells. The sporozoites emerge from a
105 host cell, begin to feed and develop into large trophozoites (Rueckert and Leander, 2008).

106 Many gregarines do not exhibit intracellular stages and are mostly epicellular parasites. The
107 gregarine life cycle typically only consists of gametogony and sporogony and only a few species
108 exhibit merogony. The sporozoites will generally develop into large trophozoites and attach to the

109 host cell with a specialised attachment apparatus (epimerite, mucron, modified protomerite)
110 (MacMillan, 1973). These specialized structures are derived from the conoid at the apical end. This
111 attachment to the host cell also functions in feeding in that the cytoplasm of the host is taken up by
112 the attached parasite (i.e., myzocytosis) (Valigurová et al., 2007). Two mature trophozoites
113 eventually pair up in a process called syzygy and develop into gamonts. The orientation of gamonts
114 during syzygy differs depending on the species (e.g. side-to-side and head-to-tail). A gametocyst
115 wall forms around each pair of gamonts, which then begins to divide into hundreds of gametes
116 (gametogeny). Pairs of gametes fuse and form zygotes, each of which becomes surrounded by an
117 oocyst wall. Within the oocyst, meiosis occurs to yield four or more spindle-shaped sporozoites
118 (sporogony). Hundreds of oocysts accumulate within each gametocyst, and are usually released via
119 host faeces or via host death and decay (Vivier and Desportes, 1990; Kuriyama et al., 2005;
120 Rueckert and Leander, 2008).

121 The gregarines are thought to be the earliest lineage of apicomplexans (Rueckert and
122 Leander, 2008) and were previously subdivided into three orders; Archigregarinida, Eugregarinida
123 and Neogregarinida (Adl et al., 2012; Grassé 1953). However, the taxonomy has recently been
124 revised (Cavalier-Smith, 2014), on the basis that it was phylogenetically unsound (Rueckert et al.,
125 2011). In this new classification, the class name Gregarinomorpha has been adopted to broadly
126 refer to all its members (i.e. gregarines, *Cryptosporidium* and Histogregarina) (Cavalier-Smith, 2014).
127 Within the various subclasses of Gregarinomorpha are Cryptogregarina, discussed above, and
128 Orthogregarinia (comprising the orders Vermigregarida and Arthrogregarida), for gregarines most
129 closely related to *Cryptosporidium* (Cavalier-Smith, 2014).

130

131 **3. Key similarities between gregarines and *Cryptosporidium***

132

133 Similarities between *Cryptosporidium* and gregarine parasites are outlined in Table 1. Key
134 similarities include the ability of *Cryptosporidium* to complete its life cycle in the absence of a host,
135 the presence of large extracellular gamont stages, syzygy (end to end pairing for reproduction) and
136 ability to adapt to their environment by changing their cell structure depending on the surrounding
137 environment.

138

139 3.1 Ability to complete its life cycle in the absence of host cells

140

141 Until recently, it was thought that *Cryptosporidium* were obligate intracellular parasites that
142 completed their life-cycle in an intra-cellular but extra-cytoplasmic (epicellular) location by pulling
143 the host cell membrane around it as an extracytoplasmic “parasitophorous sac/ membrane” that
144 sequestered the parasite from the intestinal lumen and the host cell’s cytoplasm (Tzipori and Ward,
145 2002; Dumenil, 2011). However, the initial description of the complete development of
146 *Cryptosporidium* in axenic culture (without attachment to host cells) by Hijjawi et al. in 2004,
147 revealed that *Cryptosporidium* is not an obligate epicellular parasite and this has been confirmed by
148 subsequent studies (Boxell et al., 2008; Hijjawi et al., 2010; Boxell, 2012; Yang et al., 2015;
149 Aldeyarbi and Karanis, 2016a; 2016b; 2016c). Studies by Aldeyarbi and Karanis have confirmed
150 the presence of all known life cycle stages and the production of both thin and thick-walled oocysts
151 by transmission electron microscopy (TEM) (Aldeyarbi and Karanis, 2016b; 2016c). Even when
152 *Cryptosporidium* is cultivated with host cells, it has been reported that as *C. parvum* progresses
153 through its life cycle, it becomes more extracellular with no evidence of attachment to cell lines
154 found (Borowski et al., 2010). Another study on quantitative PCR (qPCR) analysis of
155 *Cryptosporidium* growth in both cell culture and cell-free culture, reported that only ~ 5% of
156 parasite DNA could be found associated with host cells or bound to the plastic of the cell-free
157 cultures, and that the majority of parasite DNA was present in the cell culture medium (Paziewska-

158 Harris et al., 2016). These findings support the earlier observations by Pohlenz et al. (1978) and
159 Beyer et al. (2000), where intact various developing stages of parasites that are not enclosed within
160 parasitophorous sacs were found free in the calves' lumens or deep within free macrophages. This
161 is despite the fact that the very process of taking sections of intestine and processing for histology
162 analysis is likely to wash away anything not directly attached to enterocytes.

163 The ability of *Cryptosporidium* to complete its life cycle extracellularly also further
164 confirms its relationship with gregarines. For example, coelomic gregarines can also survive
165 extracellularly, even without attaching to the host intestine (Desportes and Schrével, 2013). It
166 therefore appears that *Cryptosporidium* is capable of both epicellular and extracellular
167 multiplication and development and they may both occur simultaneously in the host for mass
168 production of new oocysts (Clode et al., 2015). Preliminary work also suggests that
169 *Cryptosporidium* can complete its life cycle in water (Boxell, 2013), although this needs further
170 validation.

171

172 3.2 Extracellular gamont-like stages

173

174 The presence of gamont-like extracellular stages in the life cycle of *Cryptosporidium* was
175 first observed in a study by Hijjawi et al. (2002) and has since been reported by several
176 investigators (Hijjawi et al., 2004; Rosales et al., 2005; Karanis et al., 2008; Borowski et al., 2010;
177 Koh et al., 2013, 2014; Huang et al., 2014; Aldeyarbi and Karanis, 2016a). A previous study had
178 suggested that the presence of gamont-like stages in both cell-free and in-vitro cultures was due to
179 contaminating debris or fungal infection resembling *Bipolaris australiensis* and *Colletotrichum*
180 *acutatum* (Woods and Upton, 2007). However, TEM analysis of gamont stages (Aldeyarbi and
181 Karanis, 2016a), counters this argument.

182 Extracellular gamont-like stages have been purified from cell-free culture and *in vivo* from
183 mice infected with *C. parvum* (Hijjawi et al., 2004, Fig. 1a). Pairing of these gamont-like stages
184 with each other in a process similar to syzygy (Fig 1b), resulted in the formation of a gametocyst
185 (multi-nucleated mass)-like stage, which originated after their fusion (Fig. 1c). The identity and the
186 role of these stages are still unknown but similar cell sizes and morphologies have been observed in
187 gregarines (Leander, 2006, Leander, 2007; Alarcón et al., 2011). It has been suggested that gamont-
188 like extracellular stages might originate from sporozoites which failed to penetrate the host cells
189 and developed extracellularly into motile trophozoite stages (Hijjawi et al., 2004; Rosales et al.,
190 2005). However, given the dominance of the trophozoite stage in the life cycle (Hijjawi et al., 2004;
191 Borrowski et al., 2010; Yang et al., 2015), it is possible that they are derived from trophozoite
192 stages. Interestingly, trophozoites and developing meronts showing dividing nuclei have been
193 observed inside of unexcysted oocysts (Borrowski et al., 2010; Hijjawi et al., 2010; Aldeyarbi and
194 Karanis, (2016b) and in certain instances, *Cryptosporidium* sporozoites /zoites have the ability to
195 develop directly to sexual stages during *in vitro* cell-free culturing without appearing to go through
196 a merogonic process (Aldeyarbi and Karanis, 2016c). This plasticity in its life cycle, with the ability
197 to avoid merogony and initiate mitotic division from fused sporozoites is similar to Urosporoidea
198 (formerly eugregarines) (Rueckert et al., 2013). It is possible that gametogenesis may occur inside
199 the gametocyst-like stage and that mature oocysts are released in clumps upon its disintegration.
200 This could explain why oocysts are often seen clumped together in faecal and water samples
201 (Hijjawi, unpublished observations). Further research is required to better understand this process.

202

203 3.3 Syzygy

204

205 A defining characteristic of gregarines is syzygy (the process in which two mature
206 trophozoites pair up before the formation of a gametocyst) (Rueckert and Leander, 2008). For

207 *Cryptosporidium*, the process of syzygy (end to end pairing for reproduction) was first described by
208 Hijjawi et al. (2002). In that study, large (~10 µm) extracellular stages of *C. andersoni*, present in
209 large numbers in the faeces of infected cattle, were observed undergoing syzygy. Isolation of this
210 stage using laser microdissection and subsequent molecular characterisation confirmed that this was
211 a stage in the life cycle of *C. andersoni* (Hijjawi et al., 2002). Stages similar to these have been
212 described in the gregarine *Heliospora caprellae* (Rueckert et al., 2011). Since then, pairing of
213 *Cryptosporidium* merozoites type II/I (Borowski et al., 2010), extracellular trophozoite/gamont
214 associations (Rosales et al. 2005; Koh et al. 2014), lateral pairing between trophozoites or
215 sporozoites (Hijjawi et al., 2004; Hijjawi et al. 2010) and latero-caudal or side-by-side syzygy of
216 extracellular stages or gamonts (Aldeyarbi and Karanis, 2016a) and pairing of extracellular
217 microgametes (Aldeyarbi and Karanis, 2016c) has been reported. The latter studies by Aldeyarbi
218 and Karanis, (2016c) reported that pairing of extracellular microgametes resembles the caudo-
219 caudal syzygy of the archigregarines *Selenidiidae Selenidium pendula*, *Selenidium hollandei*
220 (Desportes and Schrével, 2013) and *Selenidium pennatum* (Kuvardina and Simdyanov, 2002). The
221 exact identity of such pairing in *Cryptosporidium* remains unknown, but it has been suggested that
222 this could be due to affinity between *Cryptosporidium* stages/gamonts rather than biological
223 purposes as in gregarines (Aldeyarbi and Karanis, 2016c).

224

225 3.4 Ability to adapt to their environment (variation in cell structure feeding modes)

226

227 Gregarines exhibit an enormous diversity in cell architecture and dimensions, depending
228 on their parasitic strategy and the surrounding environment (Leander et al. 2003b; Leander 2008;
229 Valigurová, 2012), which is also reflected in variation in feeding modes (epimerite, mucron,
230 modified protomerite) (MacMillan, 1973). This ability to adapt to their environment is also seen
231 with *Cryptosporidium*, which also appears to exhibit tremendous variety in cell structure depending

232 on the surrounding environment (Aldeyarbi and Karanis, 2016c). For example, the extension of the
233 pellicle in microgamonts may play a role the parasite's adjustment for nutrient acquisition through
234 increasing its surface area, as reported in other cell-free *Cryptosporidium* asexual and gregarine-like
235 stages (Aldeyarbi and Karanis 2016a, 2016b, 2016c). These adaptations in *Cryptosporidium* may
236 have developed to allow it to survive and grow in cell-free conditions.

237

238 **4. What does this mean for the water industry?**

239

240 The recent breakthroughs in *Cryptosporidium* biology (i.e., reclassification and confirmed
241 ability for host-free replication) could represent a paradigm shift for the water industry, particularly
242 in the area of distribution system risk. Should *Cryptosporidium* be able to access the energy
243 available in bacterial biofilms and multiply in (drinking) water distribution systems, the current
244 understanding of the impact of this pathogen and its risk assessment would need to be revised.

245

246 *4.1 Do current anti-Cryptosporidium antibodies cross react with novel gamont-like stages?*

247

248 Currently two types of antibodies are used in the water industry: (1) oocyst-specific
249 antibodies that react with the oocyst wall (various suppliers) and (2) Spor-Glo (Waterborne Inc.)
250 which is specific for sporozoites and other intracellular life cycle stages. The latter specifically
251 targets *Cryptosporidium* developmental stages by targeting antigens exposed only after excystation
252 (Boxell et al., 2008; Edwards et al., 2012; Koh et al., 2013). Non-commercial antibodies have also
253 been raised against oocysts and life-cycle stages including an antibody (anti-Cp33) that recognizes a
254 33 kDa protein on *C. parvum* but not *C. hominis* sporozoites (Jenkins et al., 2014).

255 As thick-walled *Cryptosporidium* oocysts are the infectious stage and the life form currently
256 believed to be the only stage in the life cycle capable of surviving in the environment, current

257 detection methods for *Cryptosporidium* in water have focused on detecting oocysts using the US
258 EPA 1623 method, utilising various fluorescent *Cryptosporidium* oocyst wall antibodies (DiGiorgio
259 et al., 2002). As discussed above, large gamont stages have been identified by numerous groups,
260 both *in vitro* and *in vivo*. Some studies, which have used both *Cryptosporidium* oocyst wall
261 antibodies such as Cy5-Crypt-a-Glo™ (Waterborne, Inc) and Sporo-Glo™, have reported that
262 novel gamont stages detected in biofilms did not label with either type of antibody, suggesting that
263 the surface epitopes that bind Sporo-Glo™ are not expressed by these stages (Koh et al., 2014).
264 Thus it is possible that current detection methods cannot identify all stages of *Cryptosporidium* that
265 may be present in faecal samples and/or in water supplies. However, other cell-free studies have
266 observed labelling of large gamont stages with both Sporo-Glo™ and Cryptocel (Cellabs) and these
267 stages appeared to be surrounded by a thick multi-layered wall which may be able to resist the
268 external environment and disinfection processes (Hijjawi, unpublished observations). Whether
269 current antibodies cross-react with novel gamont-like stages remains to be determined, but as
270 gamont stages are much larger than *Cryptosporidium* oocysts (10-35 µm v 5-8 µm), any observed
271 cross-reaction is likely to be dis-regarded. The fact that novel gamont-like stages have been
272 identified in faecal samples in large numbers (Hijjawi et al., 2002), suggests that this is not an
273 abnormal development as a result of a lack of a host cell, but may be a normal part of the
274 *Cryptosporidium* life cycle.

275

276 4.2 What is the susceptibility of these novel stages to disinfection?

277

278 Studies conducted to assess the survival of *Cryptosporidium* under environmental conditions
279 have focussed on oocysts only (Kothavade, 2012). The die-off rate of *Cryptosporidium* oocysts in
280 water is 0.005–0.037 10log-units day⁻¹ under natural environmental conditions (Medema et al.,

281 2006), but the fate of these novel life-cycle stages in water and the effect of disinfection procedures
282 on novel extracellular stages are unknown.

283

284 4.3 Ability of *Cryptosporidium* to survive and reproduce in biofilms

285

286 Biofilms (aggregates of micro-organisms) in both drinking water and wastewater
287 distribution networks represent a potentially significant, long-term reservoir of *Cryptosporidium*
288 because biofilm properties potentially allow them to trap and progressively concentrate
289 *Cryptosporidium* oocysts, prior to releasing them in bulk upon dislodgment of biomass (Lazarova
290 and Manem, 1995). Biofilms therefore have the potential to hold large quantities of oocysts that
291 may not be represented in water samples, potentially resulting in contamination of source waters
292 that may have been classified as oocyst free and safe for human exposure (Angles et al., 2007;
293 DiCesare et al., 2012a). Indeed, the release of *Cryptosporidium* oocysts back into the surrounding
294 water has been shown to be responsible for frequent sporadic *Cryptosporidium* outbreaks (Howe et
295 al., 2002; Wingender and Flemming, 2011). Biofilm erosion also explains the appearance of oocysts
296 in water distribution systems long after a contamination event and were suggested to be the reason
297 for ongoing recoveries of oocysts from a drinking water distribution system, following a waterborne
298 cryptosporidiosis outbreak in England (Howe et al., 2002). Biofilms have also been shown to
299 reduce solar disinfection of *Cryptosporidium* (DiCesare et al., 2012b), however, whether biofilms
300 would provide protection to *Cryptosporidium* from predation by natural zooplankton such as
301 rotifers remains to be determined.

302 As *Cryptosporidium* was believed to be an obligate intracellular parasite, it was thought to
303 lack the capacity to proliferate within biofilms and studies had focused only on the association of
304 the oocyst stage within biofilms (Keevil, 2003; Searcy et al., 2006; Wolyniak et al., 2009; Wendt et
305 al., 2015). Recently however, *Cryptosporidium* developmental stages (sporozoites, trophozoites,

306 large meronts, merozoites, microgamonts, gamont-like cells and extra-large gamont-like cells) have
307 been identified from artificial biofilms using various techniques including scanning electron
308 microscopy (SEM) and flow cytometry (Koh et al., 2013; 2014). When exposed to an artificial
309 biofilm environment *Cryptosporidium* oocysts initiated excystation and released viable sporozoites,
310 suggesting that biofilm-derived factors are therefore sufficient to initiate the excystation process
311 (Koh et al., 2014). Both asexual and sexual life cycle stages similar to the host-based life cycle were
312 identified but production of new oocysts was not reported (Koh et al., 2014). Extra-large “gigantic”
313 gamont-like cells ($30 \times 35 \mu\text{m}$) were also detected in biofilms and it is thought that they may have a
314 role in producing infective stages of *Cryptosporidium* in biofilms (Koh et al., 2014; Clode et al.,
315 2015). These very large stages have also been reported in cell-free culture and appeared to have a
316 thick outer wall (Hijjawi et al., 2002 and unpublished observations). As discussed above, it is
317 possible that mature oocysts can emerge from these gigantic stages but further research is required.

318 How *Cryptosporidium* obtains nutrients to grow in biofilms is not known but as discussed
319 above, the ability of *Cryptosporidium* to increase its surface area to optimise surface-mediated
320 nutrition (Aldeyarbi and Karanis, 2016c) may be involved. Surface nutrition through dispersed
321 micropores (pinocytosis) or osmotrophic nutrition via hair-like microvilli has been proposed in
322 gregarine species (Warner, 1968; Prensier et al., 2008), and it is possible that similar mechanisms
323 are also present in *Cryptosporidium*. However, whether biofilms or water will have the necessary
324 precursors that *Cryptosporidium* normally need to scavenge from the host, because they lack the
325 relevant biosynthetic pathways (Abrahamsen et al., 2004), is another important issue that needs to
326 be investigated.

327
328 *4.4 Implication for modelling the fate and transport of Cryptosporidium*

329

330 A number of variables can be considered for modelling the fate and transport of
331 *Cryptosporidium* within aquatic domains (Walker and Stedinger, 1999). So far algorithms have
332 been successfully applied to only one *Cryptosporidium* life stage, with pre-determined physical–
333 chemical– and biological–characteristics: the thick-walled oocyst. This has been assumed to have
334 specific coefficient parameters, for instance, for buoyancy, weight, size, settling velocity, sensitivity
335 to UV light and chlorine, biological activity (i.e., ability to replicate) etc. Clearly, the validity of the
336 numerical models and algorithms currently available must be revised on the basis of potentially
337 different properties, peculiar to the novel life stages.

338

339 **5 Research needs**

340

341 In addition to the industry research needs outlined below, basic research to better understand
342 the extracellular novel life cycle stages is needed. The novel stages have already been examined by
343 light microscopy (Hijjawi et al., 2002; Hijjawi et al., 2004; Rosales et al., 2005; Koh et al., 2014),
344 immunolabelling, confocal and scanning electron microscopy (SEM) (Edwards et al., 2012; Koh et
345 al., 2014), transmission electron microscopy (TEM) (Rosales et al., 2005; Aldeyarbi and Karanis,
346 2016a) and laser microdissection coupled with PCR and sequencing (Hijjawi et al., 2002). Novel
347 gamont stages have also been purified *in vivo* from mice infected with *C. parvum* and cattle
348 naturally infected with *C. andersoni* (Hijjawi et al., 2002; Hijjawi et al., 2004). It is still unclear
349 however, where in the life cycle these stages appear, but it has been suggested that they derive from
350 trophozoites, with “a possible role in the generation of more trophozoites or merozoites for new
351 oocyst production in host cell-free and aquatic environments” (Clode et al., 2015). The abundance
352 of novel stages in faecal samples and their infectious potential *in vitro* and *in vivo* also needs to be
353 understood as well as whether the novel stages are found in all species or only in some. If it
354 transpires that novel stages are shed in low abundance and are not infectious *in vitro* or *in vivo*, then

355 some of the more sophisticated experiments outlined below may not be required. However as noted
356 previously, novel stages have been reported in large numbers in cattle faeces (Hijjawi et al., 2002).

357

358 *5.1 Disinfection studies*

359

360 Data on the survival of life cycle stages other than oocysts under the conditions they are
361 exposed to in the natural environment are required to establish the risk posed by these stages. As a
362 key barrier in drinking water supply, the ability of chlorine and chloramines, as well as other
363 disinfectants used by the water industry, to disinfect the various *Cryptosporidium* life cycle stages
364 including novel stages needs to be determined in order to fully assess water quality risks. It is also
365 important to note that not all countries (including some European Union countries) use chemical
366 disinfection or have a disinfectant residual for drinking water supplies. If *Cryptosporidium* can
367 replicate in environmental biofilms, then systems without a residual would be at greater risk.

368 Experimental viability studies of *Cryptosporidium* oocysts in water have also found that
369 oocysts were still able to transform into trophozoite forms following UV disinfection treatment,
370 although they were not able to develop any further (Belosevic et al., 2001).

371

372 *5.2 Improvements to the cell-free culture model*

373

374 Studies have reported that *Cryptosporidium* can complete its life cycle and produce new
375 oocysts in 5-8 days post-culture inoculation (Hijjawi et al., 2004; Aldeyarbi and Karanis, 2016c).
376 However, currently only a five-fold amplification of the parasite in cell-free culture has been
377 reported (Zhang et al., 2009; Hijjawi et al., 2010). This low level of proliferation is also seen in cell
378 cultures conducted with host cells, where it has been shown that host cell detachment and apoptosis
379 are major contributing factors (Widmer et al., 2000a). As cell-free culture is not limited by host-cell

380 apoptosis, improvements to cell-free culture may result in higher oocyst outputs. Other factors that
381 affect the development and proliferation of *Cryptosporidium* in host cell culture include the
382 excystation protocol, age and strain of the parasite, stage and size of inoculum and culture
383 conditions such as pH, medium supplements and atmosphere (Hijjawi, 2010; Karanis and Aldeyarbi,
384 2011; King et al., 2011; 2015). It is likely that many of the same factors will affect the proliferation
385 of *Cryptosporidium* in cell-free media. A recent study by Edwinston et al., (2016), suggested that
386 glycoproteins and free Gal/GalNAc facilitated the switch from invasive *Cryptosporidium*
387 sporozoites to replicative trophozoites and therefore increasing the concentration of Gal/GalNAc in
388 the culture medium may enhance *Cryptosporidium* proliferation in *in vitro* culture.

389 Recently, a continuous cell culture system, based hollow fiber technology, has been
390 developed that can generate very large quantities of oocysts (1×10^8 oocysts $\text{ml}^{-1} \text{day}^{-1}$) (Moranda
391 *et al.* 2016). The hollow fiber cartridge is composed of 200 μm diameter polysulfone hollow fibers
392 with a 20 kDa molecular weight cut-off (FiberCell Systems, Inc., Frederick, MD, USA). Hollow
393 fiber technology provides several unique features: (i) a large surface area for metabolite and gas
394 exchange, which are needed for efficient growth of host cells; (ii) the creation of a biphasic medium
395 providing an oxygen rich nutrient supply to the basal layer of the host cells, while permitting the
396 provision of an anaerobic nutrient rich supply to the apical side mimicking the gut; (iii) the ability
397 to obtain high numbers of *in vitro* cultured *C. parvum* oocysts for biochemical and molecular
398 studies (Moranda *et al.* 2016). The ability to generate the intestinal redox conditions is a critical
399 factor in the success of the method. The medium is based on a modified MEM medium plus serum
400 and additives that promote parasite growth (lipids, redox buffers and vitamins), based on recent
401 biochemical and genetic studies (Abrahamsen et al., 2004; Zhu, 2008; Zhu et al., 2010). For
402 example, *C. parvum* lacks fatty acid synthase II biosynthetic machinery, suggesting they are
403 dependent upon fatty acid salvage from the host (Zhu et al., 2010). In support of this, the yield of
404 parasites obtained from the continuous culture system by Moranda et al. (2016), was significantly

405 improved by including a lipid supplement, which included the omega-3 fatty acids, alinolenic acid,
406 eicosapentaenoic acid and docosahexaenoic acid. It would be interesting to trial this technology and
407 modified medium for cell-free culture as it may also dramatically increase the oocysts yield in cell-
408 free culture. The ability to culture *Cryptosporidium* in large quantities will be extremely useful in
409 drug assessment and in research on the evolutionary biology and invasion mechanisms of
410 *Cryptosporidium*. It will also be of great benefit to the water industry. For example, to date, oocyst
411 disinfection and viability studies have concentrated on *C. parvum* and little is known about survival
412 and disinfection of *C. hominis* or other species that can infect humans as only one study on UV
413 disinfection has been conducted on *C. hominis* (Johnson et al., 2005). This is because only *C.*
414 *parvum* can be readily cultured in mice (Meloni and Thompson 1996). A gnotobiotic pig model has
415 been established for *C. hominis* (Widmer et al., 2000b; Pereira et al., 2002) and *C. hominis* is also
416 reportedly capable of infecting immuno-suppressed gerbils (Baishanbo et al., 2005); however, this
417 is beyond the facilities of most laboratories and the use of animals also has ethical implications.

418

419 5.3 Development of gamont and stage-specific antibodies

420

421 New antibodies that target more of the *Cryptosporidium* developmental stages are needed to
422 further investigate the life cycle of *Cryptosporidium*. Mono- and poly-clonal antibodies against
423 *Cryptosporidium* oocysts are available from a wide range of global suppliers. All these products
424 target one or more antigens expressed on the oocyst wall and are widely used in a variety of ways,
425 including immunofluorescence microscopy, flow cytometry (FCM), laser scanning-based systems,
426 and immunomagnetic separation (IMS methods).

427

428 Oocyst wall antigens are composed predominantly of *Cryptosporidium* oocyst wall proteins
429 (COWP1- COWP8), a family of proteins that contain polymorphic Cys-rich and His-rich repeats
(Chatterjee et al., 2010). There are also knob-like structures on the inner surface of *C. parvum*

430 oocyst walls, which cross-react with an anti-oocyst monoclonal antibody (Entrala et al., 2001). To
431 the best of our knowledge, Sporo-Glo (Waterborne Inc., USA) is the only commercially-available
432 product targeting *Cryptosporidium* antigens specific for sporozoites and other intracellular life
433 cycles stages. Cross-reactivity of the currently available commercial products, with antigens or
434 epitopes expressed on the novel life stages, has not been tested and certainly warrants further
435 investigation.

436

437 *5.3 Evaluation of the ability of Cryptosporidium to survive and propagate in biofilms*

438

439 Genomic studies have shown that *Cryptosporidium* has lost the de novo biosynthetic
440 capacity for purines, pyrimidines, and amino acids and relies solely on scavenge from the host via a
441 series of transporters (Abrahamsen et al., 2004). Previous studies by Koh et al. (2013; 2014),
442 demonstrated the attachment and development of *Cryptosporidium* in an artificial biofilm
443 (*Pseudomonas aeruginosa*), suggesting that like gregarines, *Cryptosporidium* is very plastic in its
444 response to its environment and is able to survive and grow on nutrients released by biofilms.
445 Studies are required to investigate under what conditions *Cryptosporidium* can attach to and
446 multiply in bacterial biofilms representative of biofilms found in activated sludge in water and
447 wastewater utilities. This should include identifying the conditions in biofilms that trigger oocyst
448 excystation, determining if bacteria in natural biofilms can provide *Cryptosporidium* with the
449 required purines, pyrimidines, and amino acids for survival and replication and if new thick-walled
450 environmentally robust oocysts are formed in biofilms. Additional studies may show that
451 *Cryptosporidium* development in biofilms is minimal and therefore not a significant risk, however it
452 is important that this research is conducted to better inform risk models.

453

454 **6 Conclusions**

455

456 There is now growing evidence that the gregarine *Cryptosporidium* can excyst and multiply
457 without undergoing host cell encapsulation and epicellular development, and research in this area is
458 essential for more effective catchment management. The prevalence and environmental robustness
459 of gamont stages in the environment and ability of currently used antibodies to detect gamont stages
460 needs to be explored. Similarly, whether oocysts can be produced in biofilms and their infectivity
461 also needs to be examined.

462

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466

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808 Fig. 1. (a) Nomarski interference-contrast photomicrograph of an extracellular gamont-like stage
809 purified from mice after 72 hrs infection with *C. parvum* and (b) two gamont-like stages fused
810 together with two big nuclei confirming their fusion or syzygy, (c) resulting in the formation of a
811 gametocyst (multi-nucleated mass)-like stage, which originated after their fusion. Scale bar = 5µm.
812 Images reproduced with permission from Hijjawi et al., (2004).

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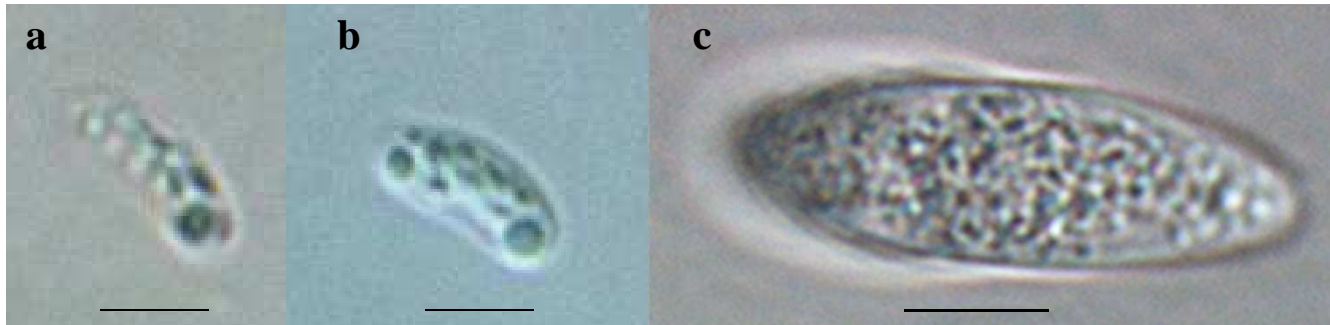
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Table 1. Similarities between *Cryptosporidium* and gregarine species.

Properties	<i>Cryptosporidium</i>	Gregarines	References
Life cycle	Monoxenous	Monoxenous	Levine, 1977; Fayer and Ungar, 1986; Rueckert et al., 2013
Location within the host cell	Occurs in the intestines enterocyte brush border	Occurs in the intestines enterocyte brush border	Levine, 1984; Fayer, 2008;
Epicellular location	Yes	In some species (Gregarinoidea)	Valigurová et al., 2007; Fayer, 2008
Feeder organelle	epimerite	mucron or epimerite	Huang et al., 2004; Valigurová et al., 2007; Borowski et al., 2008; Wisner, 2011; Aldeyarbi and Karanis, 2016a;
Myzocytosis-like feeding (cytoplasm of the host is taken up by the attached parasite)	Yes	Yes	Valigurová et al., 2007; Karanis and Aldeyarbi 2011; Aldeyarbi and Karanis, 2016a
Extracellular development	Yes	Yes	Hijjawi et al., 2002; Rosales et al., 2005; Karanis et al., 2008; Borowski et al., 2010; Koh et al., 2013, 2014; Huang et al., 2014;
Ability for intracellular or extracellular asexual replication (merogony) of trophozoites	Yes	Yes	Lange and Lord, 2012; Aldeyarbi and Karanis, 2016a; Aldeyarbi and Karanis, 2016b
Undulating epicytic-like folds covering the surface of the extracellular stages	Yes	Described for some species belonging to Terragregarina.	Lucarotti 2000; Butaeva et al. 2006; Valigurová et al., 2007; Rueckert et al. 2011; Desportes and Schrével 2013; Aldeyarbi and Karanis, 2016a
Presence of parasitiphorous sac/vacuole	Double-membrane	Multi-membranous for <i>Ditrypanocystis</i> species	Butaeva et al. 2006; Valigurová et al., 2015
Presence of Apicoplast	Absent	Mostly absent	Zhu et al., 2000; Abrahamsen et al., 2004; Cavalier-Smith, 2014
Syzygy (end to end pairing for reproduction)	Present	Present	Beams et al., 1959; Vavra and McLaughlin, 1970; Hijjawi et al., 2002; Kuvardina and Simdyanov, 2002; Toso and Omoto 2007; Borowski et al., 2010; Rueckert et al., 2011; Desportes and Schrével 2013; Aldeyarbi and Karanis, 2016a; Aldeyarbi and Karanis, 2016c
Ability of sporozoites /zoites to develop directly into sexual stages without merogony	Reported for <i>C. parvum</i>	Urosporoidea (formerly eugregarines)	Rueckert et al., 2013; Aldeyarbi and Karanis, 2016c
Auto-infective oocysts	Yes	Reported in <i>Triboliocystis garnhami</i> and <i>Farinocystis tribolii</i>	Dissanaike 1955, Žižka 1972; Fayer, 2008
No. of sporozoites / oocyst	Four	Four in many species	Kuriyama et al., 2005; Fayer, 2008, Wisner, 2011
Presence of the apical complex organelles as conoid and polar rings in	Yes	Reported in <i>Mattesia grandis</i>	Vavra and McLaughlin, 1970; Aldeyarbi and Karanis, 2016b

ACCEPTED MANUSCRIPT



Highlights

- *Cryptosporidium* now grouped with gregarine parasites
- Both epicellular and extracellular development
- Can diagnostic antibodies cross-react with novel stages?
- Susceptibility of novel stages to disinfection?
- Growth in biofilms?