

A review of the characteristics of the dinoflagellate parasite Ichthyodinium chabelardi and its potential effect on fin fish populations

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- *chabelardi* and its potential effect on fin fish populations

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24 Abstract

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26 This paper focuses on the biology and ecological impacts of Ichthyodinium chabelardi (Phylum Dinophyta, Class Syndiniophyceae, Order Syndiniales), a virulent endobiotic 27 28 parasite of yolk sacs and young larvae of many species of marine fin fish. Its infections have been observed in warm and temperate open oceanic environments and crowded 29 30 marine fish tanks. The prevalence of I. chabelardi and the range of its host fishes is not well-studied, and our understanding of its life cycle is incomplete. Here we describe what 31 is known about I. chabelardi infections in fish and we compare this with several other 32 protistan parasites of fish, including Amyloodium ocellatum, Saprolegnia parasitica, 33 Sphaerothecum destruens, and the "X-cell" clades Gadixcellia and Xcellia, all of which 34 are considered emerging generalist parasites infecting a wide variety of fin fish species. 35 36 Recent findings suggest that rising seawater temperatures might lead to higher infection rates in fishes, and we expect that these changing conditions could also expand the ranges 37 of some of these parasitic species. Thus, it is essential that the fishing industry effectively 38 39 monitor fish tanks and water in the surrounding environments for the presence of zoosporic parasites including *I. chabelardi* in order to take steps to prevent large losses in 40 41 these fisheries.

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45 Introduction

46 **Objectives of this review**

47 This review focuses on the marine dinoflagellate parasite Ichthyodinium chabelardi Hollande & J. Cachon. The genus Ichthyodinium (Class Syndiniophyceae) consists of one 48 49 species, *I. chabelardi*. This species is a virulent, generalist, endobiotic parasite of embryos and young larvae of many species of marine fin fish (Class Osteichthyes) living and 50 reproducing in the ichthyoplankton, primarily in tropical and warm temperate oceanic 51 52 ecosystems (Stratoudakis et al. 2000; Shadrin et al. 2010b). I. chabelardi has been intensively studied in only a few host fin fish species. Most of the fin fish which carry this 53 parasite have been poorly investigated, even though some of the host fish species are 54 55 economically important food resources. Because there is some evidence that the 56 prevalence of other types of parasitic infections may increase with rising seawater 57 temperatures (van West 2006; Gozlan et al. 2014; Ercan et al. 2015), it is extremely 58 important to understand their mechanisms of infectivity, potential future distributions, and begin to explore approaches to mitigation before they become an even greater threat to 59 fisheries. 60

61 Until now, no review of the literature on this parasite has been available. In contrast, several compilations have been written about a closely related host generalist parasitic 62 63 protist belonging to the Hematodinium genus (also in the Class Syndiniophyceae) (Stentiford and Shields 2005; Small and Pagenkopp 2011). The infectious agent of the 64 'bitter crab disease' (BCD) or the 'bitter crab syndrome' infects many commercially 65 66 important host species around the world including crabs, lobsters and amphipods. This 67 parasite has also been implicated in several outbreaks in fishery and aquaculture facilities leading to substantial annual economic losses (Shields 1994; Stentiford and Shields 2005; 68

Small *et al.* 2012). Given that *I. chabelardi* is also a generalist parasite infecting several
different species of fish hosts with high prevalence levels, we predict that it, too, could
have a significant impact on commercial fin fish populations.

72 In this review we firstly discuss the phylogeny, biology and ecology of *I. chabelardi* focusing on its morphology and life cycle and the pathology of its infections in host 73 74 fishes. We then describe the putative geographical expansion and host shift of this emerging disease and its potential ecological roles in marine food webs. We briefly 75 76 compare the biology of *I. chabelardi* with four other emerging zoosporic parasites which commonly infect fin fish: Amyloodium ocellatum, Saprolegnia parasitica, Sphaerothecum 77 destruens and "X-cell" parasites. Finally, we appraise the future possibility of large losses 78 in production of marine fin fish to epidemics in fish farms unless measures are 79 80 implemented to prevent the spread of parasitic infections.

81

82 Phylogenetic position of Ichthyodinium chabelardi

Baldauf (2003, 2008) divided all eukaryotic organisms into eight supergroups based upon 83 consensus phylogeny. The supergroup Alveolata consists of a very diverse range of 84 marine and freshwater protists belonging to three lineages: dinoflagellates, apicomplexans 85 and ciliates. Presently the dinoflagellates (Phylum Dinozoa) include the following large 86 clades based on both molecular and morphological characteristics: Phylum Perkinsozoa 87 (Norén et al. 1999), Class Noctiluciphyceae (Gómez et al. 2010), Class Dinophyceae 88 89 (core dinoflagellates) (Pascher 1914), and Class Syndiniophyceae (to which 90 Ichthyodinium chabelardi belongs) (Loeblich III 1976). There are also several small clades, one of which includes the genus Oxyrrhis (Lowe et al. 2010). 91

92	Within the alveolates there are phototrophic, heterotrophic, and mixotrophic species. Most
93	of the core dinoflagellates are thought to be photosynthetic (including mixotrophs), but
94	there are also some heterotrophic species. Approximately half of all known dinoflagellate
95	species in all classes taken as a group are heterotrophic (Lessard and Swift 1986). Some
96	species of heterotrophic dinoflagellates contain plastids that lack a functional
97	photosynthetic apparatus (Robledo et al. 2011). The Phylum Perkinsozoa, Class
98	Syndiniophyceae, Class Blastodinophyceae, Class Noctiluciphyceae and the Oxyrrhis
99	clade are all groups of heterotrophic protists which lack functional chloroplasts and have
100	diverged into clades that are currently considered separate from the Class Dinophyceae
101	(core dinoflagellates) (Gómez et al. 2010; Bachvaroff et al. 2014).
102	<i>I. chabelardi</i> belongs to Class Syndiniophyceae which is composed of diverse lineages of
103	mainly endobiotic parasites, with several clades consisting entirely of uncultured species
104	(Moon-Van der Staay et al. 2001; Guillou et al. 2008). The Syndiniales are exclusive to
105	marine environments. Because of their endoparasitic lifestyle, they have unusual
106	adaptations, making them almost unrecognizable as dinoflagellates when inside their
107	hosts. The Syndiniales include at least five major marine alveolate (MALV) groups,
108	typically referred to as MALV I, II, III, IV, and V (but sometimes interchangeably as
109	Syndiniales I – V) (Guillou et al. 2008; Chambouvet et al. 2011). MALV II is currently
110	recognized as the most diverse clade and contains some of the more thoroughly studied
111	species, including numerous members of the genus Amoebophrya. Most members of
112	MALV II have been identified as parasites of dinoflagellates, although some also infect
113	radiolarians. Members of MALV IV infect other metazoans including Hematodinium, and
114	Syndinium, two well-characterized parasites of crustaceans. MALV III and MALV V
115	currently consist entirely of environmental sequences, but form well-supported clades

(Guillou et al. 2008). Finally, MALV I includes Ichthyodinium spp. as well as members 116 that infect ciliates (Euduboscquellidae) (Coats 1988; Coats and Heisler 1989; Harada et 117 al. 2007; Jung et al. 2016). A schematic representation of the position of MALV I relative 118 119 to other members of Syndiniales and Dinophyceae is provided in Figure 1A. Ichthyodinium chabelardi was originally described as a syndinid dinoflagellate by 120 121 Hollande and Cachon (1952) with additional observations on its morphology and biology (Hollande and Cachon, 1953). Morphologically, I. chabelardi shares key features with 122 other dinoflagellates. These include trichocysts, condensed chromatin in the nucleus of 123 124 trophic stages, a dinospore (or zoospore) with two lateral flagella, and an alveolate pellicle 125 synthesized inside alveolar membranes (Hollande and Cachon 1952; Gestal et al. 2006; 126 Simdyanov et al. 2016).

127

The phylogenetic placement of *I. chabelardi* within the Alveolata superphylum remains 128 controversial. Gestal et al. in 2006 using molecular methods placed this parasite at the 129 base of the Perkinsozoa lineage, renaming the genus Perkinsoide (Gestal et al. 2006). 130 However, analysis of the SSU and LSU regions show poor support for this change, and 131 132 zoospore ultrastructure is more similar to other groups of dinoflagellates than the 133 Perkinsozoa. As with recent authors (Mori et al. 2007; Skovgaard et al. 2009; Simdyanov 134 et al. 2016), we retain the placement of I. chabelardi in the Syndiniophyceae until 135 additional work is done on its taxonomic affinities.

Although only one species has been formally described thus far, genetic analysis of the
SSU rDNA of the parasitic protists infecting a mix fish larvae of eight different species
reveal slight differences in the sequences, which suggests that there could be at least two
different ribotypes (98% of identity - 22 nt/ 868bp) in the sample (Sørensen *et al.* 2014).

These form separate clades within MALV I – 3 when aligned with other environmental
SSU rDNA sequences (Figure 1B). These two ribotypes have been detected to date in
distinct oceanic regions (Figure 2). These results provide the evidence that this group
might be more complex than previously described and composed of genetically distinct
parasites, as has been characterized for other syndinean parasites such as *Amoebophrya*spp. (Guillou *et al.* 2008).

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148 Life cycle of *I. chabelardi*

Ichthyodinium chabelardi was originally described as an infectious agent of sardines from
the Mediterranean Sea (Alger Bay) by Hollande and Cachon (1952), who made additional

151 observations on its morphology and biology (Hollande and Cachon 1953). These

descriptions reveal a complex life cycle that might differ depending on the parasite and/or

153 host species, with infections in some hosts appearing to have two generations of schizonts

154 while others have been observed with three (Figure 3). Subsequently, free-swimming

155 zoospores are released which can then infect new hosts.

156 The first evidence of infection appears in the yolk sac of embryos after gastrulation

157 (stages VI to XI, Dulčić 1998; Meneses et al. 2003). Infected eggs have no sign of

158 penetration on their surface, suggesting that the parasite may pass through the hole for

sperm penetration before the hole is completely closed (Yuasa *et al.* 2007).

160 Three successive stages of schizonts have been described by Hollande and Cachon (1953):

161 **Primordial schizonts**: The smallest stages of the parasite appear to be unicellular spheres

162 (maximum 1 to 3 per egg) with a diameter of approximately 8-15 μ m, less than 20 μ m

163 (Hollande and Cachon 1952, 1953; Meneses et al. 2003; Yuasa et al. 2007; Sorensen et al. 2014). These trophocytes absorb the vitellus material of the host in a central vacuole and 164 remain uni-nucleated, although mitotic divisions following by transversal divisions may 165 166 occur at this stage. Then, the nucleus undergoes multiple mitotic divisions without subsequent cytoplasmic division and the primordial schizont rapidly becomes a large 167 multinucleated structure of 100-140 µm. The cytoplasm forms cylindrical projections 168 169 around each nucleus and each of these units become a secondary schizont ($\sim 20-30 \mu m$) 170 that eventually separates from the rest in a budlike manner.

Secondary schizonts: These are cylindrical or in a racket-like shape, about 20 µm in length (Hollande and Cachon 1953). Lecithin starts to concentrate in these forms. This early secondary schizont first grows, then begins to divide by longitudinal bipartitions. As the posterior poles of the two daughter cells remain attached, the parasites assume a triangular form, resembling to a 'rosace'. Secondary schizonts can then form a long cord (up to 1–2 mm) in layers of successive groups of eight cells connected by their poles (Hollande and Cachon 1953).

Last generation of schizonts: Oblong (cylindrical) schizonts are released from the cord, 178 179 which become spherical after a series of divisions. At this phase, the yolk sac is opaque and entirely occupied by uni-nucleated parasitic cells. This last generation of schizonts 180 produce zoosporangia, which are released outside the host, in the water, generally after 181 hatching. The yolk sac breaks causing the death of the newly hatched larvae. In some 182 larvae, the parasite burst occurs immediately after hatching, while it takes more than 10 h 183 after hatching in others (Mori et al. 2007). The number of sporangia released from a yolk-184 sac larva of the yellowfin tuna *Thunnus albacares* has been estimated to be about 4×10^4 185 cells (Yuasa et al. 2007). 186

187 In the water, the sporangia undergo one to two divisions and become flagellated (Hollande and Cachon 1953). These spores have trichocysts (Gestal et al. 2006), which are known to 188 be involved in the host attachment in other Syndiniales (Miller et al. 2012). Nuclei of 189 190 released spores exhibit numerous nuclear pores and have chromatin masses, permanently condensed in circular or oval profiles of DNA. In some cases, structures like rhoptries and 191 microtubule-organizing centres, similar to specific apicomplexan characteristics, have 192 193 been observed at this developmental phase (Gestal et al. 2006). These structures may have similar function during the host invasion as the dense bodies and striated strips observed 194 195 in Amoebophrya (Miller et al. 2012). Spores remain actively swimming for a few days, 196 but cannot survive in a free-living state beyond that (Hollande and Cachon 1953). Spores 197 of different sizes (Skovgaard et al. 2009) seem to correspond to different generations of 198 division (Shadrin et al. 2015). The fate of the parasite outside of the host is not known at present. 199

200 Variations observed from this complex life cycle: The three generations of schizonts, first described on sardines from the Mediterranean Sea, were also observed in the 201 202 mackerel off Portugal (Meneses et al. 2003). However, secondary schizonts seem to be 203 absent on sardines from Portugal (Borges et al. 1996; Gestal et al. 2006) and several other 204 host species, such as Plectropomus leopardus from Japan (Mori et al. 2007) and Thunnus 205 albacore (Yuasa et al. 2007). In this case, primary schizonts lead directly to the last 206 generation schizonts, without producing the rosace and cord. No zoosporangium was formed in Plectropomus leopardus from Japan (Mori et al. 2007); instead, schizonts 207 208 themselves transformed into zoospores within 10 minutes after release.

209

210 Host range and geographical distribution

211 Parasitic species appear to be highly ubiquitous in the world's oceans. Known parasites comprised more than half of both the richness and abundance of sequences within the 212 pico-nanoplankton size fraction collected in the TARA Oceans expedition, which sampled 213 214 at 68 stations across the world's oceans (de Vargas et al. 2015). Within the TARA study, the Syndiniales featured prominently and over 2000 species-level OTUs were assigned to 215 the MALV I clade (which includes Ichthyodinium spp.). Additionally, parasitic 216 217 interactions were the most common type of interaction identified from network analysis of this global sequence dataset (Lima-Mendez et al. 2015). 218

219 There is evidence that parasitism by certain Syndiniales species could play a role in 220 regulating the populations of their hosts. This has even been suggested as a means of 221 controlling harmful algal blooms since many toxic dinoflagellates have been identified as Syndiniales hosts (Taylor 1968). In the Penzé estuary in France, Chambouvet et al. (2008) 222 consistently found successive blooms and declines of four different dinoflagellate species, 223 224 with prevalence of different Syndiniales species corresponding to or shortly lagging each 225 of the four dinoflagellate blooms. It is clear that in certain cases Syndiniales species can heavily impact the dynamics of their host populations, and given the incredible diversity 226 227 of hosts, their regulatory effects on a wide range of marine organisms, including fishes, are understudied. 228

Microscopic and molecular analysis of pelagic samples of ichthyoplankton have revealed the presence of *I. chabelardi* and relatives in the yolk sacs of fertilized eggs, embryos and larvae in various species of fish at widely distributed sites in the South China Sea (Shadrin *et al.* 2010 a and b), South Pacific (Yuasa *et al.* 2007) and in the North Eastern Atlantic and the Mediterranean (Dulčić 1998; Skovgaard *et al.* 2009). Up to now, *I. chabelardi* is reported to infect at least 13 different marine fish species including several commercially 235 important species such as *Maurolicus muelleri* (Guelin) and *Sparus aurata* L., *Trachurus*

236 trachurus L., Micromeesistius poutassou (Risso) and Engraulis encrasicolus L. and

237 Scophthalmus morhua L. (see Table 1 for more details on observations in certain host

species, Figure 2 for geographical distribution, and the Aquasymbio website:

239 http://www.aquasymbio.fr/fr).

240 The presence of this pathogen has not been reported elsewhere in the ocean in wild host

fish species but has become a pest in some fish farms (Mori *et al.* 2007; Yuasa *et al.*

242 2007). No data has been collected for the prevalence of *I. chabelardi* at most other sites at

243 which general surveys for dinoflagellates have been conducted. Most sampling sites

244 where *I. chabelardi* has been detected have been in warm water. The only surveys in cold

245 water were near Denmark (Sørensen et al. 2014). It is therefore clear that I. chabelardi

and relatives are generalist parasites infecting many bony fish species and are widely

247 distributed throughout the oceans. However, most of the studies thus far focus on few

ecosystems and broader surveys are required today to establish the global distribution of

theses parasites.

250

251 Infectivity and ecological significance

252 The prevalence of *Ichthyodinium* parasites in ichthyoplankton samples has been

253 investigated in various euphotic zones (North Atlantic, Mediterranean, South China and

254 South Pacific waters). These shorter-term studies highlight a surprisingly high prevalence

255 (Pedersen et al. 1994; Stratoudakis et al. 2000; Skovgaard et al. 2009). For example, field

studies in the North Atlantic revealed infection prevalences of 37% in sardine eggs in

257 1999 and 46% in mackerel eggs in 2000 (Stratoudakis *et al.* 2000; Meneses *et al.* 2003).

In the South Pacific the prevalence was nearly 100% for yellow fin tuna in 2007 (Yuasa *et al.* 2007).

More recently, Shadrin et al. (2010b) confirm this high prevalence at Nha Trang Bay, 260 261 Vietnam between 2001 and 2010 detecting Ichthyodinium-like parasite in representatives of eight families of fish. These long-term observations reveal also an increase of parasitic 262 263 prevalence year after year leading to a substantial proportion of fish mortality at early developmental stages. Hence, infection prevalence was no more than 1% in 1993, but 264 reached up to 98% in 2006. Similarly, a recent study on the leopard coral grouper fish in 265 266 southern Japan reveal that parasitic prevalence attributed to Ichthyodinium sp. ranged in the same batch from 1.9% in 1990 to 31.8% in 1995 (Mori et al. 2007). 267 268 Shadrin et al. (2010a) have suggested that infection by I. chabelardi has been responsible 269 for the significant disappearance of populations of fish species in warm water off the coast of Vietnam. The quantitative effect of infection by this parasite elsewhere awaits further 270 271 long-term investigation. However, present data suggest that I. chabelardi is an emerging 272 pathogen and is spreading into new marine ecosystems and possibly into freshwater ecosystems (Sørensen et al. 2014). 273

Taken all together, these findings data suggest that the impact of this parasite on fish hostpopulations is still today underestimated and long-term investigation is urgently needed.

276

277 Other emerging eukaryotic parasites of fish

Ichthyodinium chabelardi is not alone in its role as an emerging threat to fisheries. Several
other protists of differing phylogenetic origins, and different life cycles and modes of

infection, also present a challenge for fishery management. Here we briefly describe themechanisms and effects of five other eukaryotic parasites.

Amyloodium ocellatum (Class Dinophyceae, Order Peridiniales) is known to infect over 282 100 species of fin fish worldwide causing marine velvet disease or amyloodiniosis 283 284 (Landsberg et al. 1994, see also the Aquasymbio website: http://www.aquasymbio.fr/fr). 285 A. ocellatum is a virulent euryhaline ectobiotic parasite that tolerates varying osmotic conditions and temperatures and is commonly observed in warm and temperate oceanic 286 environments including coral reefs and especially in crowded fish pens and aquaria (Noga 287 288 1989; Levy et al. 2007). This species can grow at temperatures of up to 40 °C (Kuperman and Matey 1999). The details of its life cycle have been worked out, and there do appear 289 to be seasonal, temperature-related dynamics of infection prevalence (Kuperman and 290 Matey 1999). Its free-swimming dinospore stage attaches to the gills and skin of hatched 291 fish (Kuperman and Matey 1999) rather than the eggs as does I. chabelardi. 292

293 Saprolegnia parasitica (Phylum Oomycota) and Sphaerothecum destruens (Phylum

294 Mesomycetozoa) are known to be significant emerging parasites and are very serious

threats for production in populations of both wild and farmed fin fish in freshwater

ecosystems (van West 2006; Rowley et al. 2013; Gozlan et al. 2014; Sarowar et al. 2014).

Both species have been spread widely by resistant carriers into new ecosystems where

these pathogens have formerly never been observed.

Several pathotypes of *Saprolegnia* can act as opportunistic and aggressive parasites of
egg, larval and adult stages of many economically important species of fish such as
salmon, trout and catfish, and are responsible for significant economic losses in salmon
hatcheries. Thoen *et al.* (2015) found many different ITS-based genotypes among 89

isolates of *Saprolegnia* recovered from water samples and egg, larval and adult salmon
tissues from 26 salmon hatcheries along the coast of Norway. A limited number of species
of *Saprolegnia* (four species) were found in tissues of salmon: *diclina, ferax, hypogyra*and *parasitica. S. diclina* (sub-clade IIIB) clearly dominated and accounted for 79% of the
recovered species. These data indicate considerable genetic variation within the
populations of *Saprolegnia* pathogens, but the relationship to pathogenicity is not known.

Sphaerothecum destruens was spread from China to Europe by a healthy freshwater
carrier species, the top-mouth gudgeon (*Pseudorasbora parva*) (Gozlan, 2014). Recently
the rate of spread within parts of Europe has been estimated to be very rapid (Ercan *et al.*2015). This pathogen is a generalist, can infect a wide range of hosts and can cause high
mortalities, especially in commercially important fish species such as Chinook salmon
(*Oncorhynchus tshawytscha*) and Atlantic salmon (*Salmo salar*).

The "X-cell" clades Gadixcellia and Xcellia (Superphylum Alveolata) are parasites 315 associated with tumor-like formations in teleost fishes, including several groups of 316 commercial interest (Freeman et al. 2017). X-cells were originally believed to be 317 318 abnormal, diseased host cells, but cells extracted from lesions in the gills and skin of various fishes reveal them to be a related group of parasitic protists. The phylogeny of 319 these parasites is newly characterized and shows an unexpected level of divergence 320 321 between the two robust x-cell clades within the Phylum Perkinsozoa (Alveolata). Both clades are basal to the Perkinsids. Fishes with x-cell infections have been observed in the 322 Atlantic, Pacific, and Southern oceans (Evans and Tupmongkol 2014; Freeman et al. 323 324 2017), but the parasites have not yet been observed, nor their sequences detected, in 325 seawater independently of their hosts. In the common dab, diseased fish had difficulty with respiration and much lower survival than healthy fish (Diamant and Vicar 1987). 326

327 It is clear that *Ichthyodinium chabelardi* is not the only infectious disease threatening the future survival of important fisheries. Many diseases have been described. Moreover, 328 eukaryotic parasites have been under-studied and neglected so far despite their major role 329 330 in fish disease. Some methods have been proposed to control the spread of these parasites in controlled environments including reduced salt concentration and various chemicals, 331 though some of these (like malachite green) have been banned due to their toxicity to 332 humans (van West 2006). Further knowledge of each of these parasites, especially their 333 early detection and predicted responses to changing ocean conditions, will be of crucial 334 335 importance to the future of fish production.

336

337 Future perspectives and conclusions

338 There are plans in the future to build new hatcheries for production of marine fin fish 339 species such as cod along the coasts where appropriate. Countries such as Norway, Scotland, Canada and southern Australia currently have only salmon hatcheries filled with 340 341 cold freshwater. These facilities could be easily modified for marine fish using cold 342 seawater. However, a dramatic increase of disease caused by Saprolegnia in freshwater 343 fish tanks has already been observed (van West 2006). This appears to be a result of two factors: the mean temperature of the ocean is predicted to increase significantly with 344 345 global warming, and furthermore the use of malachite green in the tanks to prevent the growth of parasites has been banned internationally because of its toxicity to humans (van 346 347 West 2006).

348 It is often concluded that geographical range and prevalence of emerging infectious
349 diseases has increased tremendously over the last 20 years (Harvell 1999, 2002; Sutherst

350 2001). Climate change, habitat destruction, and introduction of exotic species (hosts and parasites) have been blamed for this increase. Actually, it has recently been observed that 351 the combination of introduction of diseased fish with the rise of seawater temperature has 352 353 significantly increased the prevalence of diseases of fin fish in general (van West 2006; Gozlan 2014; Ercan et al. 2015). We therefore predict that the ocean temperature increase 354 will provide more favourable environmental conditions for parasite growth and will 355 356 inexorably lead to geographical expansion of infectious agent populations like *I*. chabelardi and other protozoan parasites. This movement could impact population 357 358 dynamics and may drive a decrease of or even cause extinction of commercially important 359 fish population stock. It is not known whether fish could have a designated area to provide protection from infection by this parasite, but cold water in the oceans near the poles 360 361 could be a reservoir if there is sufficient oxygen present.

362 It is essential that the fishing industry effectively monitor fish tanks and water in the surrounding environments for the presence of parasites and takes steps to prevent large 363 losses in fish production. At present efforts to measure the prevalence of these parasites 364 appear to be inadequate. Molecular methods may provide tools for early detection (for 365 366 example, PCR of environmental samples using Ichthyodinium-specific primers), but 367 further research is required to develop new procedures for monitoring and controlling 368 infections. Mori et al. (2007) were able to prevent horizontal transfer of infection by using UV-sterilized or ozone-treated seawater. Methods such as these might warrant further 369 370 exploration for fish tank settings.

Pathogenic species such as *I. chabelardi* and the others discussed in this paper infect a
wide variety of host species that may favor their spread. Generalist parasites have been
responsible for huge losses in populations of plants and animals in agriculture,

aquaculture and the environment (Phillips *et al.* 2008). All of the species discussed in this review are examples of emerging pathogens and equally important to the economy of the aquaculture industry (Stentiford *et al.* 2012) in a world in which food production must be increased to keep up with human population growth. They are all spread easily and rapidly by highly infective free-living stage the zoospores from infected to uninfected susceptible host species of fish and could cause epidemics especially in highly dense host populations such as in fish farms.

381

382 Figure Legends

383

Figure 1. (A) Generalized schematic of Dinophyceae phylogeny with designation of the 384 five major clades within Syndiniales. MALV I, to which Ichthyodinium spp. belong, is 385 highlighted. Adapted from Guillou et al. 2008. (B) Partial SSU rDNA phylogeny 386 (Maximum Likelihood) of Ichthyodinium spp. together with closely related environmental 387 sequences (MALV I clade 3, following the nomenclature by Guillou et al. 2008), showing 388 the two clusters described within the genus. Environmental sequences are labeled by their 389 GenBank accession number following by the name of the clone. Sequences were retrieved 390 391 from GenBank using BLAST and aligned with referenced *Ichthyodinium spp.* sequences 392 using the online version of MAFFT version 7 (https://mafft.cbrc.jp/alignment/server/), and considering the secondary structure of RNA. RAxML (Randomized Axelerated Maximum 393 394 Likelihood) was performed using the GTRCAT substitution model, using a random number seeds, and 100 replicates for the bootstrap analyses. 395

396

397	Figure 2. Known global distributions and hosts of the two established ribotypes of
398	Ichthyodinium sp.
399	

400 Figure 3. Life cycle of *I. chabelardi* as observed in mackerel eggs (from Meneses et al.
401 2003).

402

Table 1. Some examples of pathotypes of *Ichthyodinium chabelardi* reported as parasites
of fish species. Recently identities have been confirmed by both light microscopic studies

405 and molecular methods.

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415 **Conflict of Interest**

416 The authors declare no conflict of interest.

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