Continuous *in vitro* culture of the carpet shell clam *Tapes decussatus* protozoan parasite *Perkinsus atlanticus*

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ABSTRACT: Continuous *in vitro* cultures of the clam *Tapes decussatus* parasite *Perkinsus atlanticus* were established from infected gill fragments, infected haemolymph and parasite hypnospores isolated from infected gill fragments following incubation in Ray’s fluid thioglycollate medium (RFTM). No continuous cultures could be initiated from *P. atlanticus* zoospores. Cultures initiated from hypnospores yielded the highest percentage of continuous cultures (100%, 6/6), followed by cultures initiated from gill fragments (93%, 43/46) and from haemolymph (30%, 3/10). Failures to establish continuous cultures were due to microbial contamination. The source of parasite influenced the success rate, the time taken to establish cultures and the size of cultured cells. *In vitro* proliferation of parasite cells was mainly by vegetative multiplication. Zoosporulation, yielding motile biflagellated zoospores, was observed at a low frequency (<1% of dividing cells) in every culture. Morphology of cultured cells examined with light and transmission electron microscopy corresponded to that of *P. atlanticus* found in clam tissues. Cultured cells enlarged in RFTM and stained blue-black with Lugol’s solution, which are characteristics of the *Perkinsus* species cells. DNA sequences of the internal transcribed spacer (ITS) region of the ribosomal RNA gene complex matched those of *P. atlanticus*. All cultures were established in a medium designated JL-ODRP-2A that was similar in composition to the culture medium JL-ODRP-1 originally used to propagate *Perkinsus marinus in vitro*. Proliferation of *P. atlanticus in vitro* could be supported by the commercial culture medium (1:2 v/v) DME:Ham’s F-12 with fetuin.

KEY WORDS: *Perkinsus atlanticus* · *Tapes decussatus* · *in vitro* culture · Clam parasite · Ribosomal RNA gene complex · ITS · Ultrastructure

INTRODUCTION

*Perkinsus atlanticus* has been associated with mass mortalities of the carpet shell clam *Tapes decussatus* in estuarine regions of the Portuguese and Spanish coasts (Ruano & Cachola 1986, Azevedo 1989, Santmartí et al. 1995). This protozoan parasite was designated as a new *Perkinsus* species by Azevedo (1989) based on gross morphological and ultrastructural features of various stages of its life cycle. Further evidence that *P. atlanticus* belongs to the genus *Perkinsus* was provided from nucleotide sequence analysis of the small subunit (SSU) rRNA gene and the internal transcribed

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spacer (ITS) region of the rRNA gene complex (Goggin 1994, de la Herrán et al. 2000, Robledo et al. 2000, Casas et al. 2002). Recently, Ordás & Figueras (1998) reported the in vitro culture of *P. atlanticus* from the haemolymph of an infected carpet shell clam. Identification of the cultured cells was based solely on their morphology (light and electron microscopy). Sequence analysis of the SSU small subunit rRNA gene, however, showed that those cultured cells did not correspond to a *Perkinsus* organism (Figueras et al. 2000). Continuous cultures of *P. atlanticus*, therefore, still need to be established.

The development of continuous cultures of *Perkinsus* species can lead to a better understanding of this group of parasites (La Peyre 1996). Cultures of the eastern oyster pathogen *Perkinsus marinus*, for example, have been used in a wide range of studies to address the parasite’s environmental tolerance (Burrellon et al. 1994, Dungan & Hamilton 1995), virulence (La Peyre et al. 1995, Bushek & Allen 1996a, Volety & Chu 1997), genetic composition (Reece et al. 1997, Reece et al. 2001), metabolism (Soudant et al. 2000, Soudant & Chu 2001) and drug sensitivity (Krantz 1994, Faisal et al. 1999). Since the original culture of *P. marinus*, there have been reports of at least 2 additional *Perkinsus* species, *P. chesapeaki* and *P. andrewsi*, being propagated in vitro. McLaughlin & Faisal (1998) cultured 2 protozoa from the gill (G117) and haemolymph (H49) of an infected softshell clam (*Mya arenaria*). The gill isolate G117 was further characterised as a new *Perkinsus* species, *P. chesapeaki* (Kotob et al. 1999, McLaughlin et al. 2000). *Perkinsus* parasites of infected Baltic clams *Macoma balthica* were cultured by Kleinschuster et al. (1994) and Coss et al. (2001a). The parasite culture by Coss et al. (2001b) was designated as a new *Perkinsus* species, *P. andrewsi*.

Continuous cultures of *Perkinsus chesapeaki* and of *P. andrewsi* were established using procedures developed to propagate *P. marinus*. These procedures varied greatly among researchers in the type of medium used, the culture conditions and the method used for the establishment of *P. marinus* primary cultures (reviewed in La Peyre 1996). *P. marinus* and *P. chesapeaki* were propagated in the culture medium JL-ODRP-1 originally formulated to resemble the composition of eastern oyster plasma (La Peyre et al. 1993, McLaughlin & Faisal 1998). Commercial media including DMEM:Ham’s F-12 and Leibowitz’s L-15 medium, with various nutritional supplements, were used to culture *P. marinus* and *P. andrewsi* (Gauthier & Vasta 1993, Kleinschuster & Swink 1993, Kleinschuster et al. 1994, Dungan & Hamilton 1995, Coss et al. 2001a). Cultures of the 3 *Perkinsus* spp. have been established from various sources including hearts, visceral ganglia and haemolymph of eastern oysters, gills and haemolymph of a softshell clam, and heart and haemolymph of baltic clams (Gauthier & Vasta 1993, Kleinschuster & Swink 1993, La Peyre et al. 1993, McLaughlin & Faisal 1998, Coss et al. 2001a). Parasite hypnospores isolated from infected eastern oyster tissues incubated in Ray’s fluid thioglycollate medium (RFTM) were used to quickly establish *P. marinus* continuous cultures from different geographical regions (La Peyre & Faisal 1995, Bushek & Allen 1996b). It is not known which source of parasites, if any, is optimal to establish continuous cultures of *Perkinsus* spp.

This article describes procedures for the continuous in vitro culture of *Perkinsus atlanticus* and provides evidence that the cultured cells are *P. atlanticus*. Four different sources of parasite from infected carpet shell clams were tested to establish *P. atlanticus* cultures. They included gills, haemolymph, hypnospores isolated from infected gill fragments after incubation in RFTM, and zoospores isolated after zoosporulation of hypnospores in seawater. To confirm that the cultured cells were *P. atlanticus*, they were characterised by ascertaining (1) their morphology at the light and electron microscopic levels, (2) their ability to enlarge in RFTM and subsequently to stain with Lugol’s iodine solution as described by Ray (1966), and (3) the nucleotide sequence of the ITS region of the rRNA gene complex. Finally, the growth rates of 9 isolates, 3 established from the gills, 3 established from haemolymph, and 3 established from hypnospores, were compared in the culture medium JL-ODRP-2A. Concurrently, proliferation of the 3 *P. atlanticus* gill isolates was also evaluated in the commercial medium DMEM:Ham’s F-12 supplemented with fetuin.

**MATERIALS AND METHODS**

**Clams.** Carpet shell clams *Tapes decussatus* (40 to 45 mm) were collected from an intertidal natural bed in the Ria de Arousa (Galicia, NW Spain) at Vilalonga in May 2000. Fifty clams were maintained for 12 d in a 75 l tank containing 0.22 µm filtered estuarine seawater at 25 ppt and 15°C. Water in the tank was aerated and recirculated through an ultraviolet steriliser to reduce the clams’ bacterial content. Twenty clams with the highest numbers of whitish pustules on their gill surfaces (a conspicuous sign of infection by *Perkinsus atlanticus*) were selected for *P. atlanticus* isolation and culture.

**Culture medium.** The growth culture medium JL-ODRP-2A was used to establish and maintain *Perkinsus atlanticus* cultures. This medium was similar in composition to the culture medium JL-ODRP-1 originally used to propagate *P. marinus in vitro* (La Peyre et
Initiation of cultures from zoospores. Decontaminated gill fragments were incubated in RFTM (Ray 1966) for 1 wk in the dark at room temperature, to induce hypnospore formation. The hypnospores were isolated following the procedure of J. I. Navas (authors’ pers. comm.), who modified previously reported methods (Perkins & Menzel 1966, Chu & Greene 1989). Briefly, the gill fragments were dissociated with 2.5 g l⁻¹ trypsin for 90 min. The hypnospores were separated from tissue debris by progressive filtration through a series of sieves of 300, 160, 100 and 20 µm. Hypnospores retained in the 20 µm sieve were rinsed 10 times in 30 ml ASW (600 × g, 10 min). They were decontaminated with two 30 min incubations in the antibiotic solution AS and rinsed 10 times in 30 ml ASW (600 × g, 10 min). The gill fragments were dissociated with 0.1 g l⁻¹ pronase in ASW for 1 h. Hypnospores freed from gill fragments were rinsed 3 times in 30 ml ASW with 0.1 g l⁻¹ mucin (600 × g, 10 min), decontaminated with two 30 min incubations in the antibiotic solution AS (600 × g, 10 min), and were rinsed again 3 times in 30 ml ASW with 0.1 g l⁻¹ mucin (600 × g, 10 min). Hypnospores were separated from tissue debris by centrifuging in ASW (50 × g, 3 min). The resulting supernatant containing the hypnospores was centrifuged twice in ASW (400 × g, 10 min) to pellet the hypnospores. Hypnospores were seeded in 24-well plates (10⁴ cells per well) in 1 ml of JL-ODRP-2A culture medium in each well, and incubated at 26°C in a humidified chamber. A total of 6 cultures was initiated from hypnospores. The progress of the cultures was checked with an inverted light microscope on Days 3 and 13 after seeding.

Initiation of cultures from hypnospores. Some gill fragments, decontaminated as described previously, were incubated in Ray’s fluid thiglycollate medium without agar (ARFTM, Sigma #A-0465), for 1 wk in the dark at room temperature (22 to 26°C), to induce hypnospore formation (Ray 1966, Nickens et al. 2000). The gill fragments were dissociated with 0.1 g l⁻¹ pronase in ASW for 1 h. Hypnospores freed from gill fragments were rinsed 3 times in 30 ml ASW with 0.1 g l⁻¹ mucin (600 × g, 10 min), decontaminated with two 30 min incubations in the antibiotic solution AS (600 × g, 10 min), and were rinsed again 3 times in 30 ml ASW with 0.1 g l⁻¹ mucin (600 × g, 10 min). Hypnospores were separated from tissue debris by centrifuging in ASW (50 × g, 3 min). The resulting supernatant containing the hypnospores was centrifuged twice in ASW (400 × g, 10 min) to pellet the hypnospores. Hypnospores were seeded in 24-well plates (10⁴ cells per well) in 1 ml of JL-ODRP-2A culture medium in each well, and incubated at 26°C in a humidified chamber. A total of 6 cultures was initiated from hypnospores. The progress of the cultures was checked with an inverted light microscope on Days 3 and 13 after seeding.

Initiation of cultures from the haemolymph. Haemolymph (ca. 750 µl) was withdrawn with a sterile 2 ml syringe and 23 gauge (G) needle from the posterior adductor muscles of the selected clams. To avoid haemocyte clumping and reduce bacterial contamination, the syringe was filled with 750 µl of chilled saline solution with antibiotics (803 mOsm kg⁻¹) consisting of (l⁻¹) 1.773 g NaSO₄, 0.930 g KCl, 21 g NaCl, 0.350 g NaHCO₃, 5 g EDTA, 0.05 g kanamycin A, 0.025 g gentamycin, and 100,000 U penicillin G in sterile tissue culture grade water. Haemolymph was centrifuged (800 × g, 10 min, 4°C) and the cell pellet was washed (800 × g, 10 min, 4°C) 3 times with a chilled saline solution with antibiotics (803 mOsm kg⁻¹) consisting of (l⁻¹) 1.773 g NaSO₄, 0.930 g KCl, 21.5 g NaCl, 0.350 g NaHCO₃, 0.05 g kanamycin A, 0.025 g gentamycin and 100,000 U penicillin G dissolved in tissue culture grade water. Haemolymph cells, consisting of clam haemocytes and parasites, were seeded in 24-well plates with 1 ml of JL-ODRP-2A culture medium in each well and incubated in a humidified chamber at 26°C. A total of 10 cultures was initiated from haemolymph. The progress of the cultures was checked with an inverted light microscope on Days 3, 4, 12, and 22 after seeding.

Initiation of cultures from the gills. After withdrawing haemolymph, 5 mm² fragments of gills with whitish pustules were excised with sterile scissors in a laminar flow hood and added to 50 ml test tubes with 30 ml of 25 ppt artificial seawater (ASW, Hawaii Marine Imports). The gill fragments were rinsed 10 times in 30 ml of ASW. They were decontaminated with two 30 min incubations in an antibiotic solution (AS) consisting of (l⁻¹) 400,000 U penicillin G, 0.4 g streptomycin sulphate, 0.2 g gentamycin, 0.4 g kanamycin A, 0.2 mg neomycin, 0.2 g polymyxin B, and 0.4 g erithromycin in sterile ASW. Gill fragments were rinsed again 10 times in 30 ml ASW. At least 4 gill fragments were randomly chosen from each clam. Two fragments were placed directly in a 25 cm² culture flask and 2 fragments were first finely minced with a sterile razor blade before placement in a 25 cm² culture flask. All of these gill fragments were incubated in 5 ml of JL-ODRP-2A medium per flask, at 26°C. A total of 46 cultures was initiated from gill fragments of 10 clams. The progress of the cultures was checked with an inverted light microscope on Days 4, 12, and 22 after seeding.
with an inverted light microscope on Days 7, 14, 21, and 45 after seeding.

**Maintenance of cultures.** Subculturing was performed 1 mo after the cultures were started, when parasite proliferation had given rise to a high number of cells. Clam cells did not multiply in the culture medium and eventually died. Cells in culture flasks and wells were harvested and passed 3 times through a 23 G needle with a 10 ml syringe to break up cell aggregates. The cells were rinsed 3 times in culture medium (800 × g, 15 min, 25°C). Cell viability was assessed by staining with 50 mg 1⁻¹ neutral red, and 5 × 10⁶ cells were transferred into a new flask in 5 ml of JL-ODRP-2A culture medium. Subculturing was repeated every 4 to 6 wk.

**Characterisation of the cultured cells: microscopy.** The morphology of the cultured cells following subculturing was observed with a light microscope using differential interference contrast optics. The ultrastructure of 3 *Perkinsus atlanticus* isolates established from gill fragments was examined on Days 2, 14 and 90 after subculturing. The ultrastructure of 1 *P. atlanticus* isolate established from haemolymph was also examined on Day 5 after subculturing. Cells were harvested and pelleted by centrifugation at 800 × g for 15 min. Cell pellets were fixed for 2 h with 2.5% glutaraldehyde in 0.2 M sodium cacodylate buffer, pH 7.2 at 4°C. The pellets were washed twice in cacodylate buffer and post-fixed for 2 h in 2% OsO₄ in cacodylate buffer at 4°C. The cells were enrobed with 1.5% agar solution. The agar blocks were dehydrated in an ethanol series and embedded in Epon. Ultra-thin sections were double-stained with uranyl acetate and lead citrate and examined with a JEOL 100 CXII transmission electron microscope (TEM) operated at 60 kV.

**Characterisation: DNA sequencing.** Genomic DNA was isolated from cultured cells as previously described by Reece et al. (1997) for *Perkinsus marinus* cultures. Isolations were done from 5 different isolate cultures established from gills of 5 infected clams as well as from 2 haemolymph and 2 hypnospore cultures. PCR primers used were designed in a previous study to specifically target the ITS region of the ribosomal RNA gene unit of all *Perkinsus* species except *P. qugwadi* (Casas et al. 2002). The primer pair designated ‘D’ was used to amplify an approximately 675 bp fragment of the ITS region from genomic DNA of the cultured *P. atlanticus*. PCR amplification, cloning and sequencing was done as previously described (Casas et al. 2002).

The resulting ITS region sequences were subjected to BLAST searches (Altschul et al. 1990) of the National Center for Biotechnology Information (NCBI) GenBank database and were aligned to available ITS sequences for *Perkinsus* spp. (Goggin 1994—GenBank accession numbers PAU07697, PMU07700, POU07701, PSU07698, PSU07699, Robledo et al. 2000—accession number AF140295, Coss et al. 2001b—accession numbers AF252288, AF102171, S.I. Kotob et al. unpubl.—accession numbers AF091541, AF091542, AF126022, AF150988, AF150989, AF150990, G.D. Brown et al. unpubl.—accession numbers AF149876, AF150985, AF150986, AF150987, C.A. Coss et al. unpubl.—accession number AF252288, Casas et al. 2002—accession numbers AF369967, AF369968, AF369969, AF369970, AF369971, AF369972, AF369973, AF369974, AF369975, AF369976, AF369977, AF369978, AF369979). Sequences were aligned using the CLUSTAL-W algorithm (Thompson et al. 1994) in the MacVector 7.0 DNA Sequence Analysis Software package (Oxford Molecular). Parsimony bootstrap analysis of *Perkinsus* ITS sequences was performed using PAUP* 4b8.0 (Swofford 2001) with 100 bootstrap resamplings of 100 random addition replicates. Nineteen of the ITS region sequences from the *P. atlanticus* cultures were deposited in GenBank under the following accession numbers: AF441207-AF441218 and AF472517-AF472523.

**Characterisation: enlargement of cultured cells in RFTM.** The enlargement of cultured cells in RFTM was tested for 9 different isolates, 3 established from gill fragments, 3 established from haemolymph and 3 established from hypnospores. Cells of each isolate were washed and suspended in ASW, and seeded in 24-well plates with 2 ml of RFTM in each well at a density of 0.5 × 10⁵ cells per ml. The plates were incubated for 1 wk at room temperature in the dark. The diameter of 50 random cells from each well was measured before and after incubation in RFTM. Cells incubated in RFTM for 1 wk were stained with diluted Lugol’s solution (Ray 1966). Cell viability was determined with 50 mg 1⁻¹ neutral red before and after incubation in RFTM.

**Growth measurements of Perkinsus atlanticus in JL-ODRP-2A and DMEM/F-12.** The growth rates of 9 different *P. atlanticus* isolates, 3 established from gills, 3 established from haemolymph and 3 established from hypnospores, were compared in the culture medium JL-ODRP-2A. The isolates were subcultured twice before measuring their growth rates. New subcultures were performed from each flask by transferring 12 × 10⁶ cells into a flask with 12 ml of JL-ODRP-2A culture medium. Additionally, 3 culture flasks containing 3 *P. atlanticus* isolates established from gills were set up in 1:2 DMEM/Ham’s F-12 culture medium (790 mOsm kg⁻¹, pH 6.6) supplemented with 1.7 mg 1⁻¹ fetaun, 100 U ml⁻¹ of penicillin G and streptomycin sulphate, and buffered with 50 mM Hapes and 3.7 mM sodium bicarbonate (Gauthier et al. 1995). Samples of 0.5 ml were collected on Days 2, 4, 6, 9, 12, 15, 18, 22,
cells per mother cell were log10-transformed before PLSD test. Cell diameter and number of daughter cells between culture origins were tested by means of 1-way analysis of variance, using MINITAB software. Paired comparisons were performed using the Fisher’s (PLSD) test. Cell diameter and number of daughter cells per mother cell were log10-transformed before statistical analysis to assure homogeneity of variance. Differences in cell diameter before and after incubation in RFTM were tested in the same way.

**Statistical analysis.** Differences in doubling time, cell diameter and number of daughter cells per mother cell between culture origins were tested by means of 1-way analysis of variance, using MINITAB software. Paired comparisons were performed using the Fisher’s (PLSD) test. Cell diameter and number of daughter cells per mother cell were log10-transformed before statistical analysis to assure homogeneity of variance. Differences in cell diameter before and after incubation in RFTM were tested in the same way.

**RESULTS**

**Establishment of cultures in vitro**

Thirty percent (3/10) of cultures initiated from haemolymph yielded continuous cultures. Sixty percent (6/10) of haemolymph cultures were contaminated by bacteria and thraustochytrid-like organisms by Day 3. No parasite cells were observed in the remaining 10% (1/10) of the cultures after 22 d. Haemocytes were dominant and few parasite cells were seen in the wells on Day 4. On Day 12, about 35% of the parasite cells were involved in vegetative multiplication. The percentage of hypnospores in the wells was lower than 1% and zoosporulation was rarely observed. On Day 22 more than 90% of parasite cells were in vegetative multiplication.

The percentage of cultures initiated from gill fragments yielding continuous cultures was 93% (43/46). Bacteria and/or thraustochytrid-like organisms contaminated the remaining cultures. Gill fragments in culture medium gradually lost their architecture and *Perkinsus atlanticus* cells progressively became free of the tissues. Very few cells of *P. atlanticus*, free of gill tissue, were seen in the flasks on Days 4 and 8 after seeding. On Day 12, the number of free parasite cells was still low although some stages of vegetative multiplication were observed in most flasks. The percentage of hypnospores in the flasks was lower than 1% and zoosporulation was rarely seen. On Day 22, more than 90% of parasite cells were in vegetative multiplication.

In addition, some immature trophozoites arising from multiplication were observed. No difference was detected between using infected gill fragments or finely minced infected gill as inoculum.

Every culture (6/6) initiated from hypnospores yielded a continuous culture. On Day 3, 96.2 ± 0.92% of hypnospores were not dividing, 2.8 ± 1.36% were in zoosporulation and 1 ± 0.45% were in vegetative multiplication (N = 3 cultures, mean ± SD). On Day 13, more than 90% of parasite cells were in vegetative multiplication.

No continuous cultures were successfully established from zoospores. There were no signs of division or zoospore transformation in any of the zoospore cultures even though none of these cultures was contaminated. The number of live zoospores decreased progressively until no viable zoospores were observed on Day 45 after seeding.

**Morphology of cultured cells**

Cultured cells observed by light microscopy were mostly spherical, with the cytoplasm enclosing 1 or more refractile bodies (presumably lipid droplets) and a prominent vacuole (Fig. 1), which occasionally contained a vacuoplast. Nuclei were peripheral. Cell diameter varied from 2 to 37 µm. Cells usually occurred in clumps. Cell enlargement was followed by division after the first signs of cell cleavage were observed (Fig. 2). Subsequently, daughter cells became distinguishable forming tight clusters (Fig. 3). The daughter cells enlarged inside the mother cell (Figs. 4 & 5) and were freed following rupture of the mother cell wall (Fig. 6). The number of daughter cells produced by each mother cell varied from 2 to ca. 60. Counting daughter cells was difficult due to their tight clumping. The initial diameter of daughter cells varied from 2 to 5 µm. Daughter cells still in contact could divide further, thus producing large clusters of daughter cells. Cells maintained in culture without a change of medium for an extended period of time (5 mo) developed a large vacuole and could lose their lipid droplets (Fig. 7). Hypnospores with a thick wall, discharge tube, and a variable number of prezoo- spores/zoospores inside (Fig. 8) were observed in all the cultures at a low frequency. Motile biflagellated zoospores were eventually liberated from these hypnospores.

Ultrastructurally, the cytoplasm of cultured cells 2 d after subculture showed numerous vesicles, lipid droplets, endoplasmic reticulum, small mitochondria with tubular cristae and vacuoles. Some of the vacuo- ules contained vacuoplasts. Each cell nucleus showed a prominent nucleolus (Fig. 9). Losomes were also
observed between the cell wall and plasmalemma of some cells. Cells sampled 5 d after subculture had enlarged and their cell walls had become thicker (Fig. 10). Vegetative division involved fragmentation of the vacuoles (Figs. 11 & 12); then multinucleated stages with incomplete cleavage of the cytoplasm were observed (Fig. 11). Subsequently, every nucleus was surrounded by a portion of cytoplasm, which was membrane-limited. Daughter cells were very tightly packed within the mother cell wall, which remained intact (Fig. 12). On Day 14, the cytoplasm of cultured cells had a large vacuole lacking vacuoplast, lipid droplets and mitochondria, and the nucleus showed an obvious nucleolus (Figs. 13 & 14). There were cells in vegetative division in which daughter cells were loosely packed (Fig. 13). After rupture of the mother cell wall, clusters of daughter cells in close contact and surrounded by a broken mother cell wall were fre-
quently seen (Figs. 13 & 14). Cells cultured for 90 d without a change of media had enlarged vacuole with no vacuoplast and flattened remaining cytoplasm and nucleus (Figs. 15 & 16). Some vacuole protrusions and cytoplasmic vesicles were seen, whereas lipid droplets were absent (Fig. 15). Cytoplasm appeared granular with mitochondria showing tubular cristae, the nucleus was elongated and lomosomes were observed between the cell wall and plasmatic membrane (Figs. 15 & 16).

DNA sequencing

ITS region sequences for 19 different DNA clones were obtained from 9 *Perkinsus atlanticus* isolate cultures. BLAST search results indicated that the closest matches of sequences from this study were to *P. atlanticus* (accession numbers PAU07697 and AF140295) and *P. olseni* (accession number POU07701) ITS region sequences. A consensus of optimal trees found from parsimony analysis of *Perkinsus* ITS region sequences is shown in Fig. 17. The numbers shown at the nodes indicate bootstrap support values. *P. atlanticus* ITS sequences from this study grouped with other *P. atlanticus* sequences and the *P. olseni* sequences with 100% bootstrap support in a clade that was sister to ITS sequences from *P. marinus*. Many of the sequences from the isolate culture DNAs were identical to *Perkinsus* sequences previously obtained by specific amplification of *Perkinsus* sequences from *P. atlanticus*-infected host tissue DNA (Casas et al. 2002).

Ray test

There was a significant (p < 0.001) increase in the diameter of cells incubated in RFTM. The mean (±SD, range) cell diameter was significantly larger in RFTM compared to control medium (Fig. 18).
The diameter of cultures established from gill fragments was 8.8 ± 2.93 µm (5 to 20 µm) before incubation in RFTM and 25.3 ± 8.08 µm (10 to 60 µm) after incubation in RFTM. The mean cell diameter of cultures established from haemolymph was 10.0 ± 5.02 µm (5 to 22.5 µm) before incubation in RFTM and 26.0 ± 9.06 µm (10 to 60 µm) after incubation in RFTM. The mean cell diameter of cultures established from hypnospores was 7.3 ± 4.41 µm (2.5 to 27.5 µm) before incubation in RFTM and 15.4 ± 4.29 µm (7.5 to 32.5 µm) after incubation in RFTM. The percent viability of cells, after incubation in RFTM, from cultures established from gill fragments was 34.2%, from haemolymph 28.3%, and from hypnospores 47.1%. All enlarged cells stained blue-black with Lugol's solution.

Quantitation of growth

A typical growth curve for cultured cells is shown in Fig. 18. During the first couple of days after seeding the flasks, cells enlarged and accumulated lipid droplets in the cytoplasm. On Day 2, there were significant differences (p < 0.0001) in mean cell diameter between cultures established from gill fragments, haemolymph and hypnospores. Mean (±SD) cell diameter for cultures established from gill fragments was 8.7 ± 3.54 µm (N = 201, range: 4.9 to 31.8 µm), from haemolymph 10.0 ± 4.56 µm (N = 216, 4.9 to 36.7 µm), and from hypnospores 7.3 ± 3.22 µm (N = 214, 2.4 to 29.4 µm). On Day 4, the mean diameters of cells from cultures established from gill fragments (10.1 ± 2.48 µm; N = 163, 5 to 15 µm) and haemolymph (10.4 ± 2.61 µm; N = 252, 5 to 20 µm) were significantly higher than the mean diameter of cells from cultures established from hypnospores (9.0 ± 2.62 µm; N = 297, 5 to 32.5 µm). Cell division usually started on Day 4. On Day 6 more than 50% of cells were dividing by vegetative multiplication. A few zoosporulating cells (less than 1% of dividing cells) were observed in every culture flask. The mean (±SD) diameter of cells involved in vegetative multiplication was 16.0 ± 3.55 µm (N = 83, 7.4 to 27.0 µm) for cultures established from gill fragments, 19.7 ± 6.04 µm (N = 44, 9.8 to 44.1 µm) for cultures established from haemolymph, and 13.0 ± 3.62 µm (N = 78, 7.3 to 31.8 µm) for cultures established from hypnospores. The mean number of daughter cells per mother cell (±SD) was 12.5 ± 5.74 (N = 78, 4 to 32 cells) for cultures established from gill fragments, 20.4 ± 14.10 (N = 36, 6 to 60 cells) for cultures established from haemolymph, and 8.7 ± 7.41 (N = 71, 2 to 60 cells) for cultures established from hypnospores. There were significant differences in mean diameter of cells in vegetative multiplication (p < 0.0001) and in the number of daughter cells per mother cell (p > 0.0001) between cultures established from different sources.

In JL-ODRP-2A, cultures established from haemolymph, gill fragments and hypnospores differed in their period of highest increase in cell number (i.e., log phase of growth). Log growth phase for cultures established from haemolymph occurred between Days 6 and 15; from gill fragments between Days 12 and 15; and from hypnospores between Days 12 and 18. Log growth phase for cultures in the commercial medium DME:Ham's F-12 supplemented with fetuin occurred between Days 12 and 18. Mean doubling times (±SD) for cultures established from haemolymph was 26.2 ± 6.89 h; from gill fragments 29.6 ± 8.61 h; and from hypnospores 42.3 ± 29.84 h. No significant difference in doubling times between cultures established from haemolymph, gill fragments and hypnospores was found (p = 0.55). Mean doubling times (±SD) for cultures established from gill fragments incubated in DME:Ham's F-12 was 84.5 ± 84.46 h. No difference in doubling times was observed between cultures established from gills in medium JL-ODRP-2A and those DME:Ham's F-12 (p = 0.32).
DISCUSSION

Continuous cultures of *Perkinsus atlanticus* were established for the first time. The morphology of cells in cultures, at the light and electron microscopy levels, were typical of *P. atlanticus* and other *Perkinsus* spp. Cultured cells incubated in RFTM produced enlarged cells (hypnospores) that stained blue-black in Lugol’s solution, a response characteristic of *Perkinsus* spp. and used in routine diagnosis. Species identification of the cultured cells was accomplished through analysis of nucleotide sequences of the ITS region of the rRNA gene complex, which matched that of *P. atlanticus*.

Cultures were successfully established from 3 sources of parasites, the gills and haemolymph of infected carpet shell clams and from hypnospores isolated from infected gill fragments after incubation in RFTM. However, no culture could be established with isolated zoospores. Cultures initiated from gill fragments and hypnospores yielded the highest percentage of continuous cultures, 93 and 100% respectively. The majority of cultures initiated from haemolymph became contaminated with bacteria or protozoa, and only 30% yielded continuous cultures. All contaminated cultures were discarded because past attempts to eliminate microbial contaminants generally have been unsuccessful and a drain on resources (J. F. La Peyre pers. obs.).

*Perkinsus* spp. cultures have been established from a variety of tissues including hearts, visceral ganglia and gills (Kleinschuster & Swink 1993, La Peyre et al. 1993, McLaughlin & Faisal 1998, Coss et al. 2001a) but there is little information on the optimal source of parasites to use. The success rate in establishing continuous cultures can be used as a criterion for selecting a tissue as a source of parasites. In this study, the success rate of establishing continuous cultures of *P. atlanticus* depended primarily on the absence of microbial contamination because all cultures were initiated from infected clams. Oyster hearts were used for the first time to establish *P. marinus* cultures because this organ is relatively free of microbial contaminants compared to other oyster tissues (La Peyre et al. 1993). In the current study, gills of carpet shell clams were used instead of hearts to establish *P. atlanticus* cultures because the clam heart is traversed by the intestine, thus increasing the chance of microbial contamination. Gills of *T. decussatus* are also one of the clams’ organs most infected with *P. atlanticus* (Azvedo 1989, Rodríguez & Navas 1995). Using gill tissues has the advantage that sites of infections can be easily located by looking for abscesses and hence tissues with a high parasite load can be selected to initiate cultures. Depuration of the clams, thorough rinsing of gill tissues in saline (i.e. 20 times) and decontamination of the gill tissues in antibiotic solutions were essential steps in reducing microbial contamination to establish continuous cultures of *P. atlanticus* from carpet shell clam gills. The 93% success rate in establishing *P. atlanticus* continuous cultures from gills of infected clams indicates the gill tissue is a good choice as a source of parasites.

Haemolymph also has been used to establish *Perkinsus* spp. cultures (Gauthier & Vasta 1993, Kleinschuster et al. 1994, Dungan & Hamilton 1995, McLaughlin & Faisal 1998). In the current study, microbial contamination of cultures initiated from haemolymph was more frequent than in cultures initiated from gills or hypnospores. This is not unexpected since bivalve haemolymph is seldom sterile and microorganisms such as algae and bacteria are often found in haemocytes following feeding. Establishing continuous cultures of *Perkinsus* spp. from haemolymph is therefore more difficult than from other tissues. Perhaps the decontamination procedure before seeding should be improved, although within haemocytes microorganisms cannot be rinsed off and the efficacy of antibiotic decontamination solutions is greatly reduced. One advantage of using haemolymph is that the mollusc does not have to be sacrificed, possibly enabling mollusc investigations on the contribution of different parasite strains during the progression of *Perkinsus* infection.

Another criterion for selecting a tissue as a source of parasites could be the time taken to establish continuous cultures. *Perkinsus marinus* cultures, for example, were established more rapidly from hypnospores than from hearts of eastern oysters (La Peyre & Faisal 1995). Culture flasks (25 cm²) seeded with $10^9$ *P. marinus* hypnospores contained up to $10^7$ cells per flask after 2 wk in cultures while 4 to 8 wk were generally needed to obtain similar numbers of *P. marinus* cells in cultures initiated from hearts of infected eastern oysters. In *P. atlanticus* cultures initiated with hypnospores, the parasite cells began to divide and produce new cells earlier than in cultures established from clam gills or haemolymph. The number of cultures that can be produced from the hypnospores isolated from a single clam can also be high because millions of hypnospores can be isolated from a heavily infected clam. Hypnospores have been used to establish a number of *Perkinsus* spp. cultures (La Peyre & Faisal 1995, Bushek & Allen 1996b, McLaughlin & Faisal 1998). However, the success rate in establishing cultures from hypnospores is generally much lower than the success rate in establishing cultures from oyster hearts because of microbial contamination. About 90% of cultures initiated from oyster heart tissues yield continuous cultures of *P. marinus* compared to about a 20 to 30% success rate for cultures initiated from *P. marinus* hypnospores (J. F. La Peyre pers. obs., D. Bushek pers. comm.). The much
higher success rate (100%) in establishing *P. atlanticus* cultures from hypnospores is probably due to the thorough decontamination procedure used in the current study. Gill fragments were decontaminated before incubation in RFTM and again after 1 wk incubation in RFTM before inoculation of the culture flasks. Isolated hypnospores are therefore another good source of parasites to establish continuous cultures of *P. atlanticus*.

No culture could be established when *Perkinsus atlanticus* zoospores were added to culture medium. It is not known whether the failure of zoospores to yield continuous cultures was related to the isolation method (centrifugation at high speed, 3000 × *g*) or just to the nature of zoospores. Results from this study contrast with reports of the transformation of zoospores into trophozoites and the subsequent growth in other *Perkinsus* spp. (Kleinschuster et al. 1994, La Peyre & Faisal 1995, Coss et al. 2001a). The role of zoospores in the life cycle of *Perkinsus* spp. is still unclear and further studies on the fate of zoospores are needed.

Continuous cultures of *Perkinsus atlanticus* can be established from a variety of parasite sources with varying levels of success. Success rates of establishing cultures were provided for the first time and will be useful to other researchers in selecting a source of parasites to establish continuous cultures of *P. atlanticus*.

Vegetative multiplication of *Perkinsus marinus* in its host occurs by successive bipartition in which karyokinesis is followed by cytokinesis (binary fission) (Perkins 1996). Light microscopy and TEM observation of *in vitro* *P. atlanticus* cultures suggested that vegetative multiplication occurs mostly by multiple fission; that is to say, multiple karyokinesis occurred before cytokinesis took place. Sunila et al. (2001) observed with TEM that schizogony was the most common mode of cell division of cultured *P. marinus* cells and that binary fission occurred rarely.

Zoosporulation yielding motile zoospores was observed at a low (<1% of dividing cells) frequency in every *Perkinsus atlanticus* culture established from gills, haemolymph and hypnospores in the culture medium JL-ODRP-2A. Zoosporulation of cultured vegetative cells of *Perkinsus* spp. established from several clam species has also been reported in culture media (Kleinschuster et al. 1994, McLaughlin & Faisal 1998, Coss et al. 2001a). In contrast, zoosporulation of vegetative cells from established cultures of *P. marinus* has not been reported. The significance of these observations to species identity and host specificity is not yet known.

A characteristic of protozoa in the genus *Perkinsus* is their ability to enlarge in RFTM and stain blue-black with Lugol’s solution. This characteristic discovered by Ray (1952) is the basis of a common diagnostic test for infection (Ray 1952, 1966, reviewed by Bushek et al. 1994). Only 1 *Perkinsus* species, *P. qugwadi* in bay scallops, does not enlarge in RFTM but there is doubt that this parasite belongs in the genus *Perkinsus*. In addition to the inability of *P. qugwadi* to enlarge in RFTM and to stain blue black with Lugol’s solution, differences in morphology and considerable differences in DNA sequences of the ITS region of the ribosomal RNA gene complex between *P. qugwadi* and other *Perkinsus* spp. have been found (Coss et al. 2001b, Casas et al. 2002). In this study, the size of
cultured cells incubated in RFTM increased significantly and the enlarged cells stained with Lugol’s solution. However, cultured cells incubated in RFTM were generally smaller than hypnospores isolated from infected gills (38.9 ± 10.4 μm, range 19.7 to 64.1 μm; S. M. Casas pers. obs.). Reduced enlargement of cultured cells compared to hypnospores in RFTM also has been observed in previous studies (Gauthier & Vasta 1993, Kleinschuster & Swink 1993, La Peyre et al. 1993). In vitro studies with cultured cells indicate that enlargement can be greatly increased by the addition of lipid and other nutrients to RFTM (La Peyre 1996, Wagner et al. 2001). The larger size of Perkinsus in gill tissues incubated in RFTM could reflect additional nutrients available from the carpet shell clam tissue.

Agreement in morphology between cultured cells and Perkinsus cells from host tissues is essential for genus identification but not sufficient for species identification. In some cases, protozoans that were initially described as Perkinsus spp. based partly on their morphologies were later found not to belong in the genus Perkinsus (McGladdery et al. 1991, Goggin et al. 1996, Ordás & Figueras 1998, Figueras et al. 2000). Currently, as mentioned above, classification of P. qug-wadi, is questionable because of noted differences in morphology from that which is the typical morphology of Perkinsus spp. (Blackbourn et al. 1998). Therefore, the morphology of putative Perkinsus spp. must be examined with extreme care at both the light and electron microscopic levels before they are included in this genus.

Other characteristics and data such as DNA sequences can be used in combination with cell morphology to identify protozoan parasites as Perkinsus spp. at the genus level and are required for accurate species identification within this genus. In this study DNA sequences of the ITS region helped to confirm the identification of cultures as P. atlanticus. The ITS region sequences from cultures established from gills and haemolymph and hypnospores but their DTs were not significantly different. The DTs of the 9 isolates were variable but comparable to the DTs reported for other Perkinsus spp. (Kleinschuster et al. 1994, Dungan & Hamilton 1995, Gauthier & Vasta 1995, La Peyre & Faisal 1996). However, strict comparison of Perkinsus spp. growth rates is not possible because of many differences between studies, such as in media, isolates, seeding densities, proliferation assays and the period of cell proliferation selected to calculate DT (La Peyre 1996).

The experience acquired with culturing Perkinsus marinus facilitated the in vitro culture of P. atlanticus and other Perkinsus spp. The optimal culture conditions for P. atlanticus will need to be determined by measuring the effects of various physical and chemical parameters such as temperature, salinity, pH, inoculum density and nutrient concentrations on the parasite growth. As with P. marinus, P. atlanticus cultures can be used in a wide range of studies to address important questions such as about the parasite’s environmental tolerance, virulence, genetic make up, metabolism and drug sensitivity. The development of continuous cultures of P. atlanticus will lead to better understanding of this important parasite of carpet shell clams.

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