

Protoceratium reticulatum (Dinophyceae) in the austral Southwestern Atlantic and the first report on YTX-production in shelf waters of Argentina



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ABSTRACT

Protoceratium reticulatum is a dinoflagellate with a life cycle that includes a motile planktonic stage and a resting cyst stage in benthic habitat, both with a wide geographical distribution, including southern South America. *P. reticulatum* produces yessotoxins (YTX) – these can be accumulated in shellfish and show potent cytotoxicity, posing a risk to human health if contaminated shellfish is consumed. YTX have been reported from coastal shellfish of many localities, but until now it was unknown if they were present in the austral Southwestern Atlantic and also if local populations of *P. reticulatum* have the ability to produce these toxins. In this study we report the presence of YTX in plankton samples and its production in culture by two *P. reticulatum* strains isolated from the San Jorge Gulf (SJG). In addition, we describe the geographical distribution and seasonal abundance of this species based on data collected over the past two decades. The YTX cell quotas calculated from net hauls (~ 10 pg cell⁻¹) are in the same range as the toxin cell quotas observed in these two isolates. The phylogenetic analysis of sequences of the hypervariable region of the large subunit (LSU) 28S rDNA showed that the two clonal strains from the SJG were part of a monophyletic clade that subdivides *P. reticulatum* into two well-supported, divergent sub-clades. The sequences of the two strains of *P. reticulatum* from the SJG fell in the same clade as the majority of sequences of *P. reticulatum*, which belong to a geographically widely distributed evolutionary clade. *P. reticulatum* was occasionally observed from about 35° S in Uruguayan shelf waters up to 53° S on the Patagonian shelf and north of Tierra del Fuego, and it was present from coastal areas up to the shelf break zone. We recorded *P. reticulatum* in plankton samples during spring, summer and autumn but invariably in low abundance (maximum: 560 cells L⁻¹). Viable cysts of the species in surface sediments also showed a wide geographical distribution. Together, the high total abundances and high relative numerical contribution to planktonic dinoflagellate assemblages near frontal areas, emphasize the necessity to pay attention to the dynamics of this species in areas of potential risk of harmful algal bloom development.

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1. Introduction

Protoceratium reticulatum (Claparède et Lachmann) Bütschli, is a dinoflagellate with a life cycle that includes a motile planktonic stage and a benthic resting cyst stage. It has a complex taxonomic

and nomenclatural history dating back to the 19th century. First described as *Peridinium reticulatum* by Claparède and Lachmann (1858–1859), it was later transferred to the genus *Protoceratium* by Bütschli (1885). After nearly a century, Reinecke (1967) described a new species, *Gonyaulax grindleyi*, apparently ignoring the work by Bütschli (1885) and other reports of *P. reticulatum* from European waters (e.g. Wołoszyńska, 1928). Most authors considered *P. reticulatum* and *G. grindleyi* as conspecific and have used either of these binomials (e.g. Steidinger and Tangen, 1996; Taylor et al.,

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2004). The morphological study of Hansen et al. (1996/1997) supported conspecificity of the two taxa based on plate tabulation analysis of samples collected at a site close to the locality from which the type species of *Protoceratium* (*P. aceros* Bergh) was first described, concluding that the name *Protoceratium reticulatum* has priority and should therefore be used. The motile stage of *P. reticulatum* has a wide geographical distribution that comprises global coastal waters including the North Sea and the Baltic Sea, and the Mexican Pacific, the Gulf of Mexico, the Atlantic coast of North America, the east coast of Russia, New Zealand and South Africa, among other areas (e.g. Braarud, 1945; Reinecke, 1967; Steidinger and Williams, 1970; Hernández-Becerril, 1988; Hansen et al., 1996/1997; MacKenzie et al., 1998; Hällfors, 2004; Kim et al., 2004; Orlova et al., 2004). In South America, the species has been recorded in coastal waters of Argentina, Chile and Brazil (see Balech, 1988; Seguel et al., 2005; Odebrecht, 2010).

Braarud (1945) was the first author who described the spiny cysts of *Protoceratium reticulatum*, formed in cultures from a clone isolated in the Oslo Fjord, Norway. Wall and Dale (1968) successfully incubated cysts from recent sediments and the germinated mobile thecae were identified as *Protoceratium reticulatum*. The incubated cysts were recorded by these authors as identical to the Pleistocene cysts that Wall (1967) identified as *Operculodinium centrocarpum*. In the original description, *O. centrocarpum* (as *Hystriosphæridium centrocarpum*) from the Miocene appeared as a species with processes of 13–18 µm length, solid, showing radial fibrils at the insertion point and apices widened with small curved tips (Deflandre and Cookson, 1955). Wall (1967) described specimens from the Pleistocene of the Caribbean Sea that assigned to *H. centrocarpum* but transferred the species to the genus *Operculodinium*. Therefore, modern and Quaternary cysts are referred to as *Operculodinium centrocarpum* (Deflandre et Cookson) *sensu* Wall, 1967. *O. centrocarpum* is known in the paleontological record from the Eocene to recent times, having been documented from numerous sites including Holocene sediments of the northern inner shelf and the Beagle Channel in Argentina (Borromei and Quattrocchio, 2001; Borel et al., 2006; Vilanova et al., 2008). It has been recorded in the plankton (Reid, 1978) and also has a wide geographical distribution in recent sediments (Marret and Zonneveld, 2003), being one of the most abundant species in some benthic dinocyst communities. It has been characterized as a cosmopolitan, temperate to tropical, neritic and oceanic species (e.g. Wall et al., 1977; Harland, 1983). In recent sediments from the Southwestern Atlantic it was first quoted by Wall et al. (1977) in a transect offshore off the Río de la Plata river, and it was also recorded from near coastal sites of the northern Argentine shelf (Grill and Guerslein, 1995; Akselman, 1999). Modern cysts show a great variability in process lengths and salinity and temperature have been shown to play an important role in this variation (Verleye et al., 2012).

A special interest in *Protoceratium reticulatum* arose toward the end of the 1990s after discovering that it produces a newly described type of toxin, the yessotoxins (YTX), responsible for shellfish toxicity (Satake et al., 1997; MacKenzie et al., 1998). Previously, it was also associated with shellfish toxicity in South African waters (Reinecke, 1967), and were suspected to be paralytic shellfish poisoning (PSP) toxins (Grindley and Nel, 1970). Yessotoxin is a lipophilic ladder shaped polyketides that was first isolated from the scallop *Patinopecten yessoensis* Jay after a red-tide in Japan by Murata et al. (1987). Since its discovery a growing number of YTX analogs (up to date almost 100) has been isolated and characterized from both dinoflagellates and shellfish (see e.g. Paz et al., 2013, and references therein). Yessotoxins are toxic to mice after intraperitoneal injection and show potent cytotoxicity against human tumor cell lines (Konishi et al., 2004). Yessotoxin in particular, is a powerful compound that increases

permeability in rat liver mitochondria (Bianchi et al., 2004). After its discovery in Japan, YTX and its congeners were also isolated from shellfish in many coastal sites (e.g., Ciminiello et al., 1997; MacKenzie et al., 1998; Arévalo et al., 2004; Aasen et al., 2005; Quilliam et al., 2006; Morton et al., 2007; Howard et al., 2008; Belin and Zouher, 2009). In South America YTX were detected in shellfish from Chile and Brazil (Yasumoto and Takizawa, 1997; Schramm et al., 2010). Production of YTX has been shown to occur during all phases of growth in batch culture of *P. reticulatum* strains and the toxins are contained within cells as well as in the culture medium (Paz et al., 2013). Besides *P. reticulatum*, two other gonyaulacoid dinoflagellate species, *Lingulodinium polyedrum* (Stein) Dodge and *Gonyaulax spinifera* (Claparède et Lachmann) Diesing, have been shown to also produce YTX (Tubaro et al., 1998; Draisci et al., 1999; Rhodes et al., 2006). *Lingulodinium polyedrum* and *Gonyaulax spinifera* are also present in the Southwest Atlantic (Balech, 1988). However, it still was unknown if YTX are present in waters of the austral Southwest Atlantic and also if *Protoceratium reticulatum* and/or *L. polyedrum* and *G. spinifera* have the ability to produce these toxins in this region.

In this study we report on the first detection of YTX in natural phytoplankton samples from the Southwest Atlantic and present YTX profiles of strains of *Protoceratium reticulatum* isolated from the San Jorge Gulf (SJG). A phylogenetic analysis of the hypervariable region of the large subunit (LSU) ribosomal gene was carried out to assess the genetic relationships between the South American isolates and other isolates. We also compiled data collected over the past two decades on the distribution and seasonal abundance of this species in the region and present information on the distribution of resting cysts in plankton samples and surface sediments collected in various sectors of the coastal shelves of the Southwest Atlantic.

2. Materials and methods

2.1. Field collection of plankton and sediments. Hydrographic data

Sampling procedures were performed on board research cruises conducted to the time-series station Estación Permanente de Estudios Ambientales – EPEA (Permanent Station of Environmental Studies, ~15 nmi off Mar del Plata, ~monthly intervals from May 1994 to April 1995), the Southern Patagonian shelf (March–April 2000 and 2004, cruises EH-03-00 and EH-03-04, respectively), the Argentine-Uruguayan Common Fishing Zone (AUCFZ) (November 2001, EH-09-01), the shelf break area and Southern Patagonian shelf (October 2005, March–April 2006, PD-GEF Patagonia 1 and 2, respectively), the Patagonian littoral and San Jorge Gulf (January 2010, OB-01-10), a sampling in the Beagle Channel, Tierra del Fuego Island (March 2012), and a cruise from the Beagle Channel to Mar del Plata (March–April 2012, PD Patagonia Austral 2012). Surface sediment sampling was also conducted at the port of Mar del Plata and the neighboring coastal localities of Chapadmalal and Miramar in January 1986 (Fig. 1, and Supplementary Material: Table 1, Fig. 1).

Supplementary Fig. S1 related to this article can be found, in the online version, at doi:10.1016/j.hal.2015.03.001.

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For quantitative analysis of phytoplankton, water samples were collected with Niskin bottles at standard depths in cruises to the EPEA station, the AUCFZ and the Southern Patagonian shelf, while three depths were sampled according to the fluorometric profile (surface, at fluorescence maximum and at a selected depth within the stratum below) in cruises to the shelf break area and Southern Patagonian shelf. These samples were preserved with Lugol's

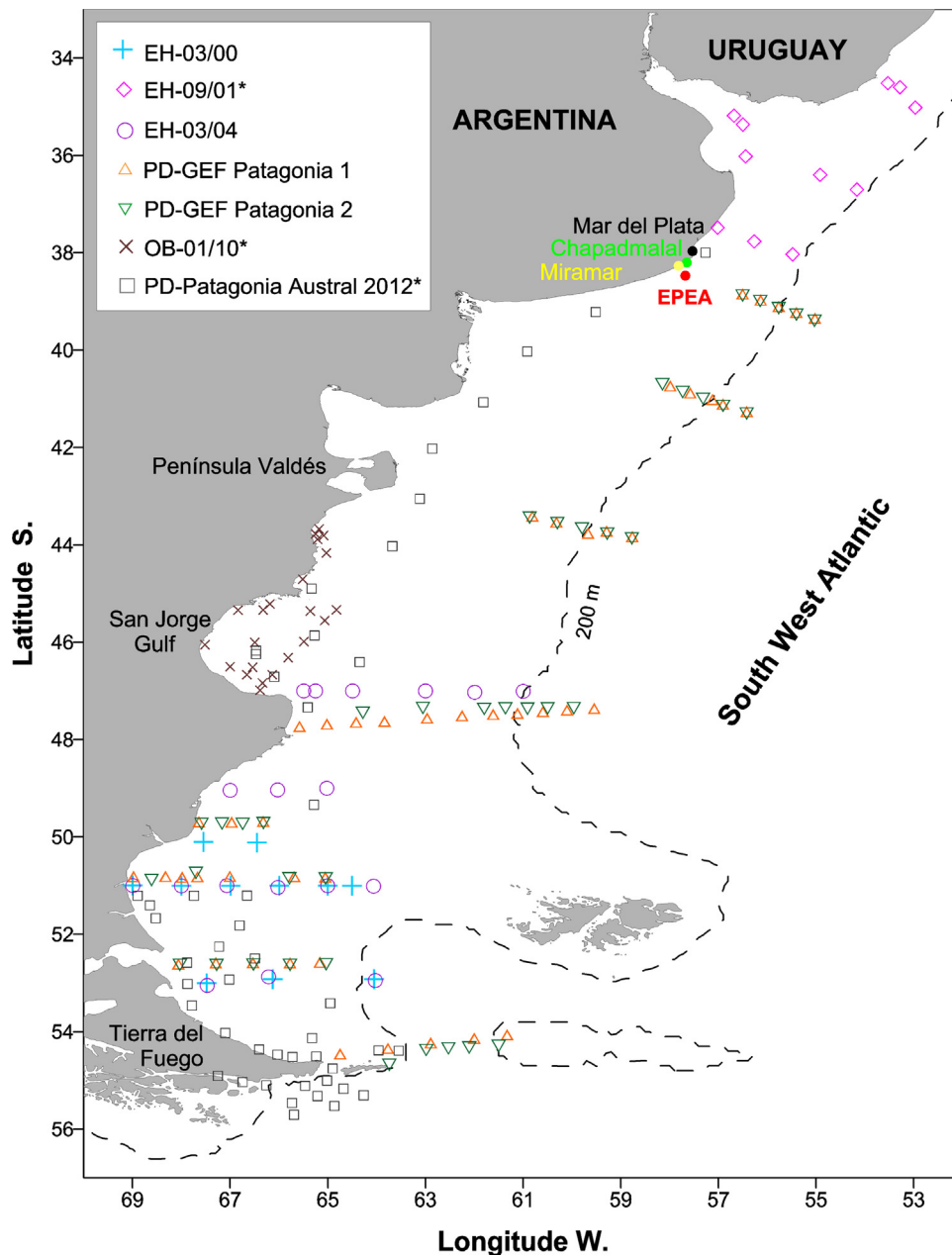


Fig. 1. Austral Southwestern Atlantic. Location of the study area showing the EPEA station (Permanent Station of Environmental Studies), sampling localities and geographical positions of stations where plankton and/or sediment samples were collected during several cruises. An asterisk in the legend indicates those cruises in which sediment samples were collected.

solution. During the cruise from the Beagle Channel to Mar del Plata, vertical net tows were conducted at each station through the upper 20 m of the water column with a 20 μm -mesh Nitex plankton net. Net hauls were adjusted to a volume of 1 L with filtered (3 μm) sea water and an aliquot of 20 mL was fixed with 0.4 mL of Lugol's solution.

Different sampling devices were used to collect sediments for dinoflagellate cyst studies: a Piccard dredge, a snapper and a manually operated grab bucket (Supplementary material Table 1). The first two centimeters were collected for analysis from the snapper and grab bucket samplers, while representative aliquots were collected from dredge samples. Surface samples were obtained from the top centimeter of box coring sediments collected during cruises to the Patagonian littoral zone and SJG, and from the Beagle Channel to Mar del Plata. Sediment samples

were kept in dark cool (4 °C) conditions to prevent cyst germination until analysis.

Temperature and salinity profiles were measured with a CTD (Seabird SBE 19).

2.2. Plankton analysis

Identification and enumeration of cells in quantitative plankton samples were done following the Utermöhl method (Hasle, 1978) with an Olympus IX70 inverted microscope coupled to an image capture system (CCD-RGB camera, Sony DXC 151A). Subsamples of vertical net tows (see Section 2.6 for plankton collection and size-fractionation) were counted in duplicates in Sedgewick-Rafter chambers, using a phase contrast LEICA DMIL LED inverted microscope. The whole chamber bottom was scanned at

200× magnification. Additional qualitative observations were done by UV epifluorescence microscopy (Leica DM 2500) of thecal plates stained with Calcofluor according to Fritz and Triemer (1985). Moreover, selected subsamples were mounted onto glass stubs and sputter coated with Au–Pd following procedures by Ferrario et al. (1995), and examined by scanning electron microscopy (Jeol JSM-6360 LV). Author names of algal species was used according to Guiry and Guiry (2014).

2.3. Dinoflagellate cyst analysis from surface sediments

Different processing methods for sediments were used to analyze dinoflagellate cyst assemblages according to the cruise under consideration.

During cruises to the EPEA station, the AUCFZ and the Patagonian littoral and SJG, an aliquot (3 cm³) of sediment from each station was sonicated for 1 min and subsequently sieved through gauze of 125 µm and 25 µm mesh size with filtered seawater. The 25–125 µm size fraction was resuspended in a known volume of filtered seawater and neutralized formaldehyde solution. A Sedgewick-Rafter chamber was used for cyst enumeration and a minimum of 250 dinocysts was counted per sample. Abundance was expressed in cysts cm⁻³ of wet sediment. Cyst assemblages processed by this method included organic-walled and calcareous cysts.

A qualitative analysis of *Protoceratium reticulatum* cysts (presence/absence) was done for samples from coastal localities. During the cruise from the Beagle Channel to Mar del Plata, an aliquot (5–10 g wet weight) of sediment from each station was sieved through 150 µm and 10 µm Nitex[®] screens, and one calibrated tablet of *Lycopodium* spores was added in each sample as exotic markers (Stockmarr, 1971) to allow calculations of concentrations in cysts g⁻¹ of wet sediment. The fraction between 10 and 150 µm was treated with cold 10% hydrochloric acid. Zinc chloride was used as heavy liquid to separate the organic fraction by density gradient. Residues were sieved and collected on a 10-µm mesh and then mounted between slide and cover-slide in glycerine jelly. Organic-walled dinoflagellate cysts (“dinocysts”) were identified using a transmitted-light microscope (Nikon Eclipse 600) at a magnification factor of 600× and 1000× and a minimum of 250 dinocysts was counted per sample. Permanent slides (LPUNS-PA) are stored at the Colección Palinológica, Laboratorio de Palinología (INGEOSUR-UNS), Bahía Blanca, Argentina.

2.4. Isolation and culturing

At station C43 (PD Patagonia Austral 2012 Cruise, see Supplementary material Table 1 for geographic position) 100 mL of sea water from 3 m depth were filled into a PE flask and kept at 4 °C until inspection. Single *Protoceratium reticulatum* cells were isolated from this sample under a stereomicroscope (Olympus SZH-ILLD) by micropipette. Cells were transferred into individual wells of 96-well tissue culture plates (TPP, Trasadingen, Switzerland) containing 300 µL of K medium (Keller et al., 1987), prepared from 0.2 µm sterile-filtered natural Antarctic seawater in 1/10 of the original concentration. Isolated cells were then incubated at 15 °C under artificial light at a photon flux density of 50 µmol photons m⁻² s⁻¹ on a 16:8 light:dark photoperiod. After 3–4 weeks, two unialgal isolates (provisionally named A2 and H1) were transferred to polystyrene cell culture flasks each containing 50 mL of ½ strength K medium and were maintained thereafter under the same conditions as described.

For toxin analysis, two cultures for each strain were grown at two different dates in 70 mL polystyrene cell culture flasks at the same environmental conditions as described before. Cell density was determined by settling lugol-fixed samples and counting

>800 cells under an inverted microscope. Dense cultures (cell concentration: 8,510 mL⁻¹ and 16,791 mL⁻¹ for A2; 10,930 mL⁻¹ and 18,909 mL⁻¹ for H1) were harvested in 50 mL Falcon tubes by centrifugation (Eppendorf 5810R, Hamburg, Germany) at 3,220 × g for 15 min. Each pellet was transferred to an Eppendorf microtube and again centrifuged (Eppendorf 5415, 16,000 × g, 5 min) and stored frozen (–20 °C) until use. Yessotoxins were analyzed separately for each pellet and cell quota for each strain was calculated as a mean of both cell pellets.

For molecular genetic analysis, one culture for each isolate was grown and harvested exactly as described before. Cell concentrations at harvest time in this case were 4,440 mL⁻¹ for strain A2 and 10,610 mL⁻¹ for strain H1, respectively. Cell pellets for sequence analysis were also stored at –20 °C until use.

2.5. Molecular phylogenetic analysis

Cell pellets were used to extract DNA with the DNeasy Plant Mini Kit (Qiagen, Hilden, Germany) according to the manufacturer's instructions and as detailed in Töbe et al. (2013). Resulting genomic DNA was used in PCR to amplify the D1/D2 hypervariable region of the large sub-unit (LSU) of the ribosomal operon with the primers D1R (forward) and D2C (reverse) (Scholin et al., 1994) and PCR chemistry and cycling conditions as in Töbe et al. (2013). Products of PCR were purified with a QIAquick PCR Purification Kit (Qiagen, Hilden, Germany) and used in Sanger sequencing with the same primers as in PCR with a BigDye[®] Terminator v3.1 Cycle Sequencing Kit (Applied Biosystems, Darmstadt, Germany). After clean-up of sequencing reaction products with a DyeEx 2.0 Spin Kit (Qiagen), the sequences were read on an ABI 3130XL Genetic Analyzer (Applied Biosystems). Primer sequences were removed from contigs after assembly of forward and reverse sequences in Geneious Pro 5.4.4 (Biomatters Ltd., Auckland, New Zealand). A BLAST search was conducted with the sequences of isolates H1 and A2 and the resultant D1/D2 region LSU sequences of other *Protoceratium reticulatum* isolates were retrieved from GenBank. Sequences of *P. reticulatum* and a set of other sequences were aligned in MAFFT v7.017 using a plugin for Geneious Pro. The additional sequences corresponded to those that were used to analyze phylogenetic relationships among gonyaulacoid dinoflagellates in Howard et al. (2009) and an additional set of sequences representing the five ribotypes of the *Alexandrium tamarense*/*A. fundyense*/*A. catenella* species complex (Supplementary material Table 2, Bolch and de Salas, 2007; Daugbjerg et al., 2000; Ellegaard et al., 2003; Herrera-Sepúlveda et al., 2013; Higman et al., 2001; Kim et al., 2006; Kim and Kim, 2007; Mertens et al., 2012; Orr et al., 2011; Riccardi et al., 2006; Ruiz Sebastián and O'Ryan, 2001; Scorzetti et al., 2009; Walsh et al., 1998). A sequence of *Prorocentrum minimum* was included as outgroup. The resultant alignment was restricted to the length of the sequences of isolates H1 and A2 and the best fitting model of nucleotide substitution was determined according to scores of the Akaike Information Criterion in jModeltest 0.1.1 (Guindon and Gascuel, 2003; Posada, 2008). A Maximum Likelihood (ML) tree was constructed using the GTR + G model of nucleotide substitution in PhyML (Guindon and Gascuel, 2003) via a plugin in Geneious Pro. Reliability of tree topology was estimated by 200 bootstrap replicates.

Supplementary Table S2 related to this article can be found, in the online version, at doi:10.1016/j.hal.2015.03.001.

2.6. Chemical analysis of yessotoxins

For plankton collection and size-fractionation, vertical net tow concentrates of stations from the PD-Patagonia Austral 2012 cruise (Supplementary material Table 1) were sequentially filtered

through Nitex® meshes of 200, 50 and 20 μm by gravity filtration and each size fraction was split into aliquots. One aliquot was used for the extraction of lipophilic toxins.

For toxin extraction from plankton, the cell pellets from the 50 to 20 μm aliquots of the plankton net tows and *Protoceratium reticulatum* cultures were harvested by centrifugation ($3,220 \times g$, 15 min at 4 °C), suspended in 500 μL methanol for lipophilic toxins, and subsequently transferred into a FastPrep tube containing 0.9 g of lysing matrix D (Thermo Savant, Illkirch, France). The samples were homogenized by reciprocal shaking at maximum speed (6.5 m s^{-1}) for 45 s in a Bio101 FastPrep instrument (Thermo Savant, Illkirch, France). After homogenization, samples were centrifuged at $16,100 \times g$ at 4 °C for 15 min. The supernatant was transferred to a spin-filter (0.45 μm pore-size, Millipore Ultrafree, Eschborn, Germany) and centrifuged for 30 s at $800 \times g$, followed by transfer to autosampler vials.

Field samples were analyzed for multiple lipophilic toxins as described in Krock et al. (2008b). Mass spectrometric analyses of *Protoceratium reticulatum* isolates for YTX were performed on a Hypersil BDS C8 column ($50 \times 2 \text{ mm}$, 3 μm , 120 Å) at a flow rate of 0.3 mL min^{-1} using an elution gradient with two eluants (A: water and B: 95% acetonitrile/methanol (1:2, v/v) and 5% water, both eluants containing 2.0 mM ammonium formate and 50 mM formic acid). Initial composition was 40% B with a linear gradient to 100% B at 6 min, isocratic 100% B until 15 min, then returning to initial conditions. Selected transitions (precursor ion > fragment ion) are given in Supplementary material Table 3. Yessotoxin was identified by its mass transitions and comparison of retention times of samples and a reference standard (IMB-NRC, Halifax, Canada). Relative abundances are based on peak area comparisons, identical response factors for all transitions are assumed.

Supplementary Table S3 related to this article can be found, in the online version, at doi:10.1016/j.hal.2015.03.001.

3. Results

3.1. Thecal and cyst morphology

Cells of *Protoceratium reticulatum* analyzed from water samples (EH-03-04 (St 256) and the EPEA station) showed morphological characteristics (Balech, 1971, as *Gonyaulax grindleyi*) which briefly were the following: subsphaeroidal with the longitudinal axis greater than the transverse, without neck and spines; epitheca shorter than hypotheca; cingulum displacement of 1–2 times its width, descendent, without overhanging; sulcus narrow, excavated; dark wall with a marked reticulated sculpture, raised ridges and a pore in each polygon; and the first apical plate with a prominent ventral pore on its right side (Fig. 2A–C). Plate formula: Po, 3', 1a, 6'', 6C, 6''', 2''', ~7S. Dimensions: 34–39 μm (length), 29–31 μm (transdiameter) ($n = 20$ cells). Its cysts (Matsuoka, 1985) were spherical, transparent, with hollow processes of variable length which were capitate or cylindrical with closed extremities; precingular archeopyle at the third paraplate, operculum free (Fig. 2D–F). Diameter (without processes): 35–45 μm ($n = 15$ cysts).

3.2. Phylogenetic analysis

The Maximum Likelihood (ML) analysis of the hypervariable region of the large subunit (LSU) 28S rDNA showed that the sequences obtained from the two clonal strains A2 and H1 from the SJG were identical to each other and to other sequences of *Protoceratium reticulatum* strains included in the multiple sequence alignment (Fig. 3). In the present ML tree, all *P. reticulatum* were part of a monophyletic clade that further subdivides into two well supported, divergent sub-clades. The sequences of the two clonal strains isolated in this study fell in the same clade as the majority of sequences of *P. reticulatum* and only three isolates of *P. reticulatum* from the west and east coasts of USA (Supplementary material Table 2) comprised the second sub-clade within *P. reticulatum*.

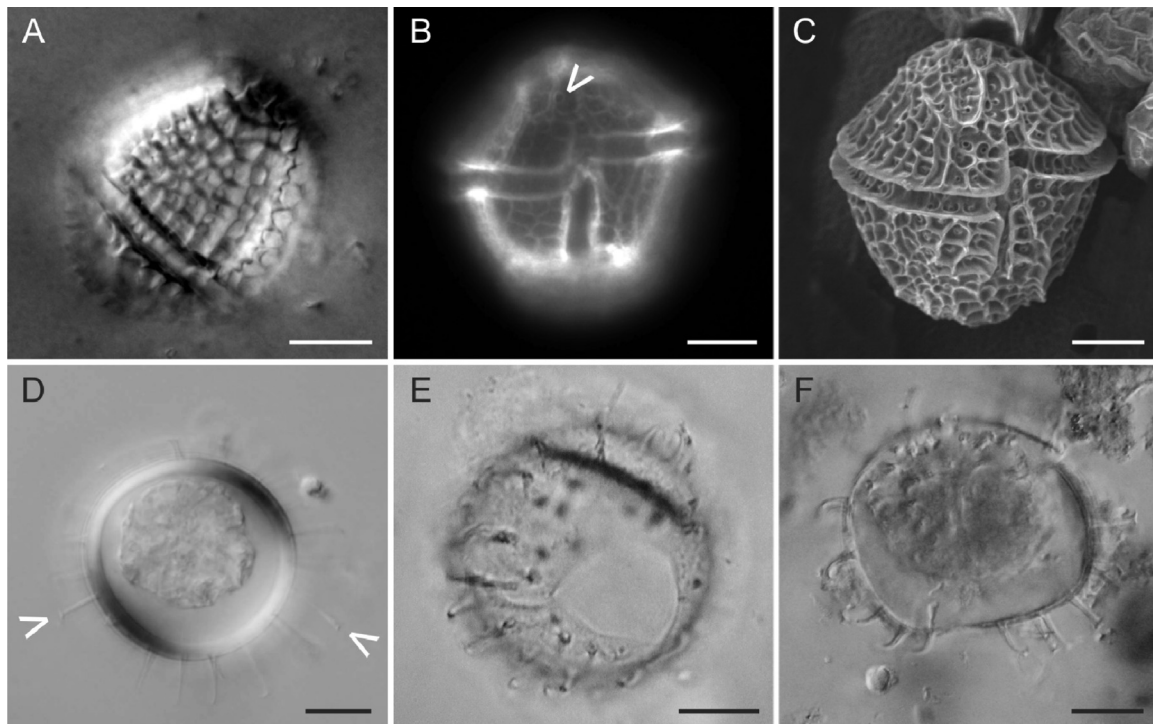


Fig. 2. Cells and resting cysts of *Protoceratium reticulatum*. Light microphotographs of motile cells in overall dorsal view showing the rough thecal ornamentation (A), and in ventral view (B, arrowhead: ventral pore on first apical plate). SEM micrograph of a theca in ventral view (C). Light microscopy views of a living cyst from a plankton sample of South Patagonia (D, arrowheads: capitate processes), and of an empty cyst showing the precingular archeopyle (E) and a cyst with cellular content (F) from recent sediments of the San Jorge Gulf. Bars: 10 μm .

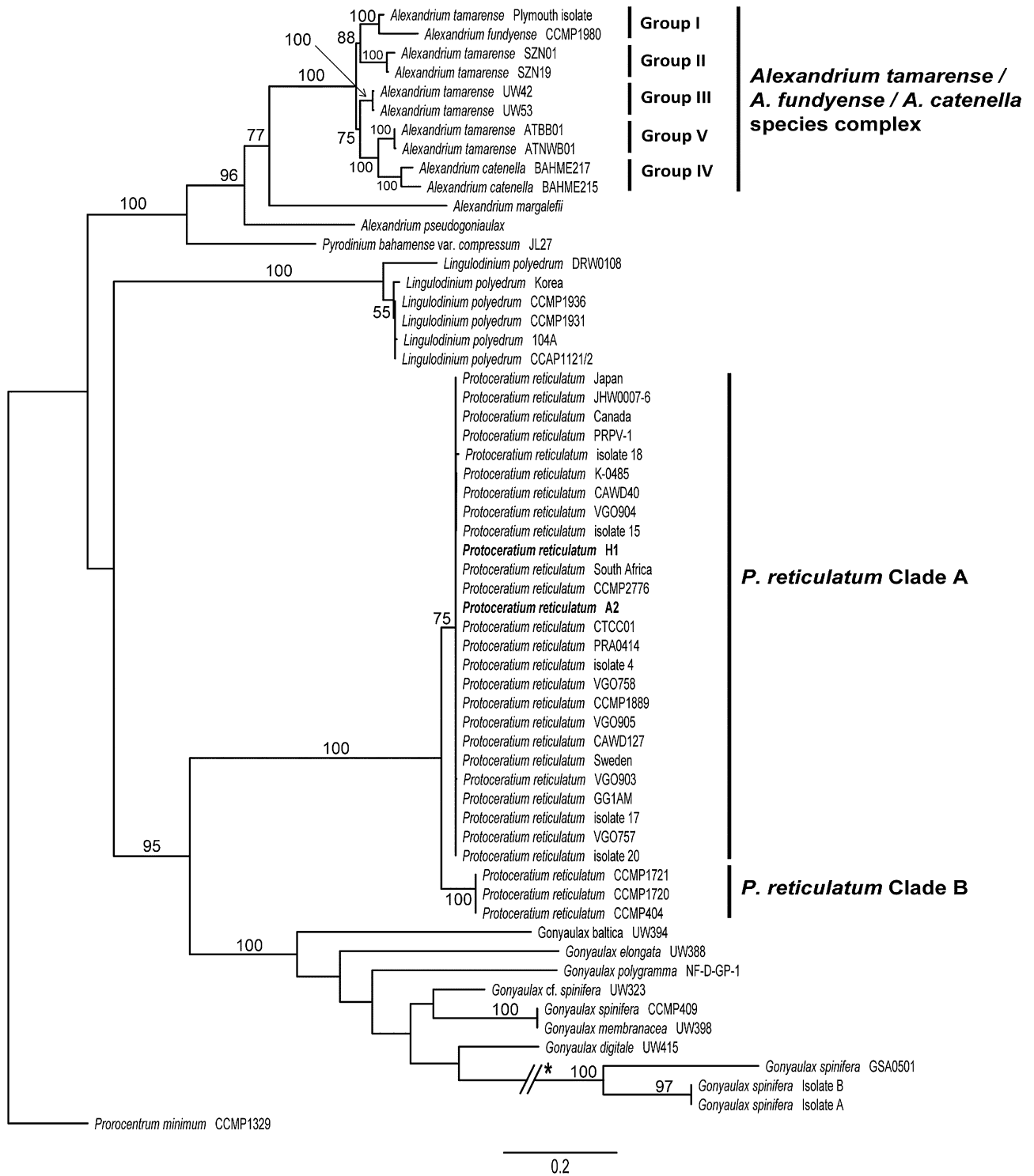


Fig. 3. Maximum Likelihood gene tree of *P. reticulatum* strains and other Gonyaulacales inferred from partial sequences of the 28S rDNA. Numbers on branches are support values >50% derived from 200 bootstrap replicates. The scale bar shows average number of substitutions per site. Sequences of the clonal strains A2 and H1 (in bold) isolated from the San Jorge Gulf fall in the same clade (Clade A) as the majority of sequences of the two divergent clades formed by strains of *P. reticulatum* in this analysis. Ribotypes/species within the *Alexandrium tamarensense*/*A. fundyense*/*A. catenella* species complex as identified by Scholin et al. (1994) and John et al. (2003) are numbered according to Lilly et al. (2007). * original branch length: 0.88.

3.3. Spatial distribution, abundance and seasonal occurrence of the motile cell stage

We found motile cells of *Protoceratium reticulatum* throughout a broad latitudinal distribution ranging from 34.52° S in Uruguayan shelf waters up to 52.63° S at the southern shelf in Argentina (Fig. 4). Geographic distribution encompassed the AUCFZ including

the mouth of the Río de la Plata up to the Magellan Strait and north of Tierra del Fuego, covering the northern Argentine shelf and central and southern Patagonia. *P. reticulatum* also showed a wide distribution across the shelf, being present at latitudes of 38–40° S from near the coast at the EPEA station (~48 m depth) up to the shelf break area (max. depth ~1529 m), and at ~45–47° S from the SJG, where it was recorded in net haul samples (PD-Patagonia

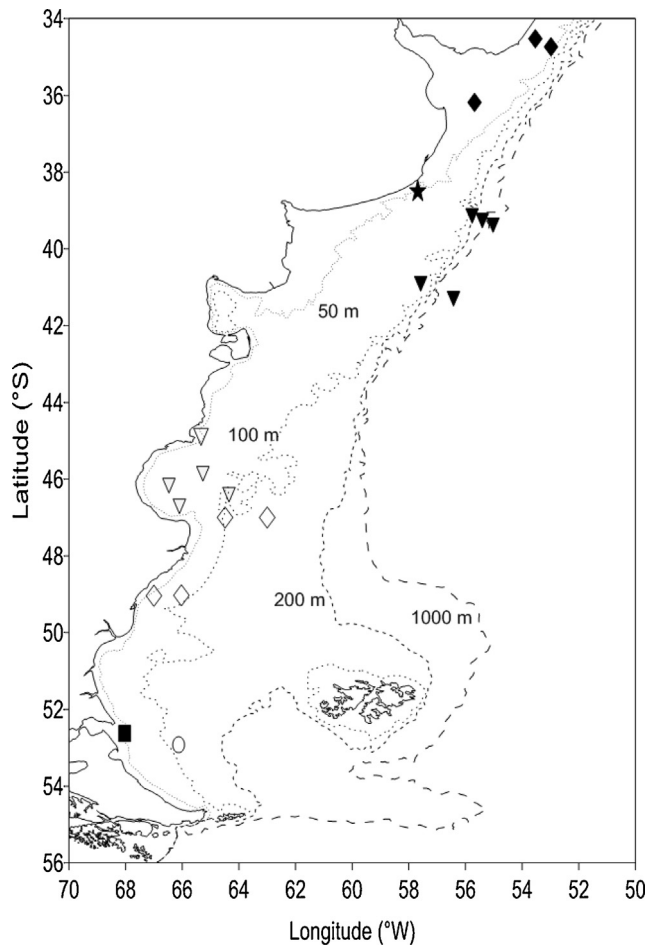


Fig. 4. Geographic distribution of the motile stage of *P. reticulatum* at the austral Southwestern Atlantic. Symbols correspond to the EPEA station (★) and analyzed cruises (○: EH-03-00; ◆: EH-09-01; ◇: EH-03-04; ▼: PD GEF Patagonia 1; ■: PD GEF Patagonia 2; ▽: PD Patagonia Austral 2012).

Austral 2012), to the central shelf. Despite its large spatial distribution, *P. reticulatum* was recorded only sporadically in quantitative samples collected during the five analyzed cruises of sufficient spatial coverage (EH-03-00, EH-09-01, EH-03-04, PD-GEF Patagonia 1 and PD-GEF Patagonia 2). Indeed, in the cruise to the AUCFZ area, *P. reticulatum* was registered at 3 out of the 12 analyzed stations, while in cruises which have covered the shelf break area between $\sim 39^\circ$ and 44° S and the Southern Patagonian shelf between 47° and 55° S, it was recorded in 5 of 30 and in 6 of 90 stations respectively. *P. reticulatum* was also detected only once during the sampling period at the EPEA station.

Abundance of *Protoceratium reticulatum* was invariably low in all the examined plankton samples, and its cells were registered in different depths of the water column. Quantitative estimations showed a maximum abundance of 560 cells L^{-1} , but values of ~ 100 cells L^{-1} were frequently observed. In some cases, although not detected in bottle samples, its cells were registered in plankton net samples.

The seasonal occurrence of the motile stage was analyzed at the EPEA station during the period from May 1994 to April 1995. Bottle samples collected at standard depths in 12 cruises showed that *Protoceratium reticulatum* appeared only once, in January 1995 (i.e. in summer at the Southern Hemisphere), and in low abundance (max. 260 cells L^{-1}) in the water column. Moreover, it was recorded only in samples collected during cruises conducted in spring (EH-09/01, PD-GEF Patagonia 1) and autumn (EH-03/00,

EH-03/04, PD-GEF Patagonia 2, PD-Patagonia Austral 2012). It should be noted that the data analyzed were from a database of species present in cruises conducted at different seasonal periods, with data selected for the occurrence of *P. reticulatum*. Both sources of information suggest that in the austral Southwest Atlantic region *P. reticulatum* cells were present in the warm period of the year.

Surface salinity and temperature values associated to the presence of *Protoceratium reticulatum* in plankton samples spanned ranges of 23–34.1 and 6.8–20.3 $^\circ C$, respectively. These values represent not only hydrographic conditions of the different water masses but also the whole period of spring, summer and autumn during which the species was recorded. The lower salinity was recorded at the mouth of the Río de la Plata, while low values were also determined north of the river in Uruguayan shelf waters (30–32.1), at the Magellan Strait (32.8–33.2), and in southern Patagonia (32.9–33.1). At the shelf break area *P. reticulatum* was present in high salinity (33.9–34.1) and low temperature waters (6.8–7.3 $^\circ C$).

3.4. Spatial distribution and abundance of resting cysts

Resting cysts of *Protoceratium reticulatum* in surface sediments were recorded in shelf areas at the AUCFZ, the EPEA station, the Patagonian littoral north of 45° S and the SJG, as well as in the three coastal localities sampled at northern Argentine shelf, spanning a latitudinal range from 34.52° S to 46.67° S and from the coast to near the 200 m isobath (Fig. 5).

At the EPEA, resting cysts with cellular content or viable cysts, as well as empty cysts of *Protoceratium reticulatum* were identified all throughout the annual period, being present in sediments from the 12 analyzed cruises (Table 1). Its abundance ranged from 8 to 470 cysts cm^{-3} wet sediment, and its percentage from 0.8% to 35% in viable dinoflagellate cyst assemblages. Viable cyst concentration were high with an average value of 94 cysts cm^{-3} wet sediment and an average percentage of 10.9% of the whole assemblage of dinoflagellate cysts. This high variability of cyst abundance at small spatial scale showed a significant negative correlation to the size grain of sediments, i.e. higher cyst concentration in sediments with smaller sediment grains (R. Akselman, unpubl. results). These results from the EPEA show that although there was a latent potential ability of benthic cysts to inoculate the water column throughout the year, detection of planktonic cells was scarce, however both in number and the period in which they were detected. In addition to *P. reticulatum*, viable cysts of *Lingulodinium polyedrum* and species of *Gonyaulax* of the *Gonyaulax spinifera* complex were also present, among other species.

Qualitatively analyzed samples at the coastal localities of Mar del Plata, Chapadmalal and Miramar showed only empty cysts of *Protoceratium reticulatum*.

Abundance of viable *Protoceratium reticulatum* cysts at the AUCFZ ranged 16–4,368 cysts cm^{-3} wet sediment with percentages of 11–83% in dinoflagellate cysts assemblages (Table 1). *P. reticulatum* cysts were recorded in sediments from the three transects, at the Uruguayan and Argentinian shelves and at the boundary line facing the Río de la Plata. High concentrations ($\geq 1,000$ cysts cm^{-3} wet sediment) and a marked variability were shown in each transect, with the highest value at the Río de la Plata (St 802). In this area and regardless of their abundance, *P. reticulatum* cysts constituted an important fraction with values generally $\geq 30\%$ of all dinoflagellate cysts. In addition to *P. reticulatum*, live cysts of *Lingulodinium polyedrum* were recorded among other species in assemblages at the AUCFZ.

At the Patagonian littoral north of 45° S and at the SJG, estimates made in samples processed without acid treatment, viable cyst concentration of *Protoceratium reticulatum* ranged 20–2,933 cysts cm^{-3} wet sediment and attained percentages of

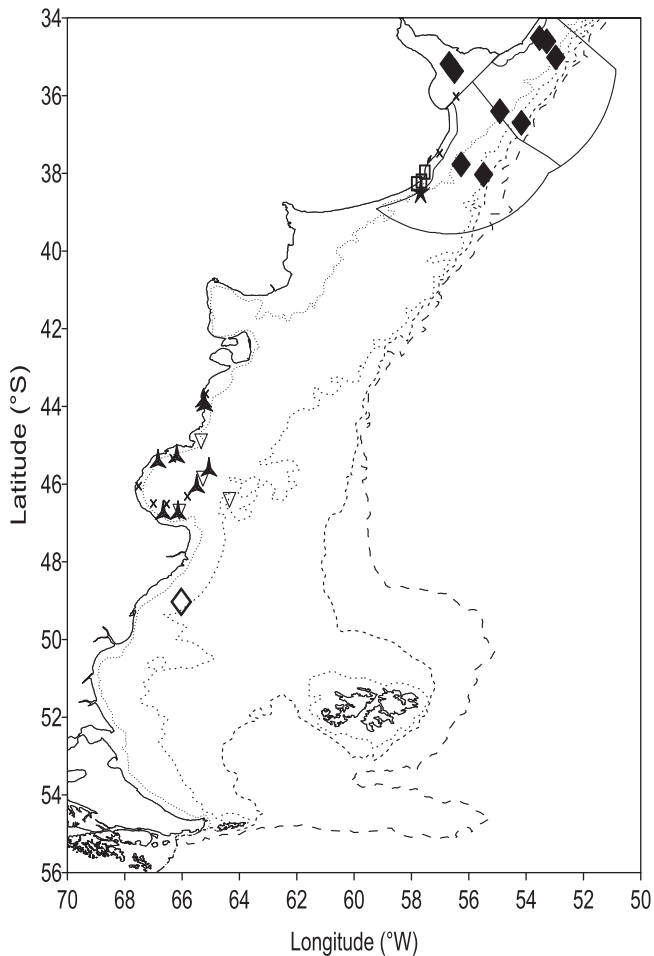


Fig. 5. Geographic distribution of *P. reticulatum* cysts at the austral Southwestern Atlantic. Symbols correspond to the EPEA station (★), the coastal localities of Mar del Plata, Chapadmalal and Miramar (□) and the analyzed cruises (◆: EH-09-01; ◇: EH-03-04; ▽: PD Patagonia Austral 2012; ▲: OB-01-10). All data points correspond to surface sediments, with exception of the EH-03-04 cruise where the record came from plankton samples. ×: stations at which *P. reticulatum* was not found.

2–33% in assemblages of calcareous and organic-walled dinoflagellate cysts (Table 1). In this area, as well as in the AUCFZ, there were locations (OB-01-10 cruise, St 2, 19) at which both cysts concentration and relative frequencies were high, suggesting the relative importance of *P. reticulatum* over others in its potential capacity to inoculate the water column. Sediment samples processed by palynological treatment, which gives information on organic walled cysts (dinocysts), showed *P. reticulatum* concentration values ranging from 66 to 401 cysts g^{-1} wet sediment (Table 1). Spatial distribution of dinocysts seemed to be heterogeneous, possibly due to the sediment grain size, as the highest concentration corresponds to muddy sediments (silty clay) and the lowest value to sites with an important fraction of sabulitic and sandy sediments (C.M. Borel, unpubl. results). Cysts of *P. reticulatum* were recorded in sediment samples from stations at which its motile cells and yessotoxins were detected simultaneously. Cysts of the *Gonyaulax spinifera* complex were dominant at the four analyzed stations from the PD-Patagonia Austral 2012 cruise (Table 1).

Moreover, viable *Protoceratium reticulatum* cysts were observed in plankton samples collected at higher latitude in south Patagonia (EH-03-04, St 256) (Figs. 2A and 5). At this position resting cysts were simultaneously observed with cells in water samples. These

Table 1

Resting cysts of *P. reticulatum* in surface sediments from cruises conducted to the EPEA station during the period May 1994–April 1995, the AUCFZ (EH-09-01) and the Patagonian shelf (OB-01-10, PD Patagonia Austral 2012). Cyst concentration in cruises to the EPEA station, EH-09-01 and OB-01-10: cysts cm^{-3} wet sediment; to the PD Patagonia Austral 2012: cysts g^{-1} wet sediment. Relative abundance in cruises EH-09-01 and OB-01-10: in assemblages of calcareous and organic-walled dinoflagellate cysts; in PD Patagonia Austral 2012: in assemblages of organic-walled dinoflagellate cysts.

| Cruise | Station | Cyst abundance | Cyst percentage |
|---------------------------|---------|----------------|-----------------|
| EH-03/94 | EPEA | 10 | 14 |
| EH-04/94 | EPEA | 60 | 7 |
| EH-06/94 | EPEA | 8 | 11 |
| EH-08/94 | EPEA | 15 | 0.8 |
| OB-06/94 | EPEA | 470 | 35 |
| OB-08/94 | EPEA | 22 | 1 |
| OB-09/94 | EPEA | 84 | 17 |
| OB-11/94 | EPEA | 14 | 2 |
| EH-01/95 | EPEA | 102 | 9 |
| EH-03/95 | EPEA | 305 | 17 |
| OB-04/95 | EPEA | 23 | 10 |
| OB-06/95 | EPEA | 14 | 7 |
| EH-09-01 | 826 | 16 | 40 |
| EH-09-01 | 825 | 80 | 11 |
| EH-09-01 | 822 | 1,120 | 37 |
| EH-09-01 | 811 | 260 | 83 |
| EH-09-01 | 809 | 346 | 33 |
| EH-09-01 | 802 | 4,368 | 26 |
| EH-09-01 | 800 | 216 | 30 |
| EH-09-01 | 792 | 1,000 | 40 |
| EH-09-01 | 795 | 57 | 50 |
| OB-01-10 | 2 | 1,467 | 24 |
| OB-01-10 | 4 | 139 | 10 |
| OB-01-10 | 9 | 106 | 2 |
| OB-01-10 | 11 | 111 | 20 |
| OB-01-10 | 19 | 2,933 | 33 |
| OB-01-10 | 29 | 889 | 10 |
| OB-01-10 | 35 | 367 | 10 |
| OB-01-10 | 36 | 20 | 2 |
| PD Patagonia Austral 2012 | C43 | 66 | 11 |
| PD Patagonia Austral 2012 | C45 | 247 | 13 |
| PD Patagonia Austral 2012 | C43N | 401 | 2.5 |
| PD Patagonia Austral 2012 | P45B | 218 | 11 |

cysts seemed of recent formation and not resuspended from sediments (i.e. they had no detritus attached), coincided with the maximum cell abundance (560 cells L^{-1}) estimated in all the analyzed cruises, and were recorded together with motile cells in all levels of the water column, observations which suggest that encystment could have been occurring.

3.5. Toxin measurements

No YTX was detected in samples from the 21 sampling points in the Beagle Channel (March 2012). Along the entire PD 2012 transect from Ushuaia to Mar del Plata, YTX was only detected at five stations in the SJG region: C43, C43N, I44, C45 and P45B (Table 2) at concentrations ranging from 24 ng net haul $^{-1}$ at station P45B up to 343 ng net haul $^{-1}$ at station I44. The yessotoxin cell quotas ranged from 4.9 to 12.7 pg cell $^{-1}$ which were calculated from the total YTX in these 20–50 μm size fractions of the net hauls and the *Protoceratium reticulatum* cell abundances in the respective samples (Table 2).

The two strains of *Protoceratium reticulatum* A2 and H1 isolated from water samples collected at station C43 had YTX cell quotas of 9.1 pg cell $^{-1}$ and 10.2 pg cell $^{-1}$, respectively. The YTX profiles of both strains were dominated by yessotoxin (>95%). The second most abundant YTX congener was probably 32-O-arabinofuranosyl-YTX (mass transition m/z 1273 > 1193) with a maximum relative abundance <5%, which has been described by Miles et al. (2005). However, an unambiguous identification of this and the following compounds was not possible due to the lack of reference

Table 2

Yessotoxin levels and *Protoceratium reticulatum* and *Gonyaulax* aff. *spinifera* cell counts per net haul (mean abundance) of the San Jorge Gulf stations (PD-Patagonia Austral 2012 cruise). Calculated cell quotas for *P. reticulatum* of these net hauls are also shown.

| Station | YTX [ng net haul ⁻¹] | <i>Protoceratium reticulatum</i> | | <i>Gonyaulax</i> aff. <i>spinifera</i> |
|---------|----------------------------------|---|---|---|
| | | Abundance [10 ³ cells net haul ⁻¹] | YTX cell quota [pg cell ⁻¹] | Abundance [10 ³ cells net haul ⁻¹] |
| C43 | 173 | 23 | 7.5 | 0 |
| I44 | 343 | 27.5 | 12.5 | 3.5 |
| C45 | 197 | 40 | 4.9 | 15.5 |
| C43N | 248 | 19.5 | 12.7 | 20 |
| P45B | 24 | 3.5 | 7.0 | 2 |

materials. In addition to the *m/z* 1273 compound, three other putative YTX could be detected at trace levels <0.5%: *m/z* 1047 >967 (41-keto-YTX), *m/z* 1157 > 1077 (OH-YTX), and *m/z* 1175 > 1095 (diOH-YTX).

4. Discussion

4.1. Taxonomy

The motile cells of *Protoceratium reticulatum* analyzed herein showed morphological characteristics and plate pattern in accordance with those described by Balech (1971, as *Gonyaulax grindleyi*) based on samples from the Southwestern Atlantic, which also correspond to the original description of *G. grindleyi* made by Reinecke (1967). In our samples from the EPEA station and South Patagonia (~10 cells of each locality were analyzed), epithelial cell configuration of the apical and intercalary plate series were 3' and 1a, and we have not observed a plate configuration of 4' and 0a. Hansen et al. (1996/1997) showed that nearly 50% of both plate configurations were present in natural samples from the type locality of *Protoceratium aceros*, the type species of *Protoceratium*, and concluded that *P. reticulatum*, *P. aceros* and *G. grindleyi* are conspecific, the first name having priority. In his monograph of the Southwest Atlantic, Balech (1988) did not consider *P. reticulatum*, although he had cited this species in two samples from the Productivity III cruise (Balech, 1971). In the latter study, he also simultaneously recorded *G. grindleyi* in the second of these samples, which was used for the first record and morphological description of this species in the Southwest Atlantic. The fact that Balech (1988) did not include *P. reticulatum*, but described *G. grindleyi* and considered *P. reticulatum* in its synonymy, could be interpreted by the application of his taxonomic criteria for defining *sensu lato* the genus *Gonyaulax*. The study of Hansen et al. (1996/1997) clarified uncertainties involved in the formal taxonomy and nomenclature regarding these taxonomic questions.

4.2. Phylogenetic analysis

The Maximum Likelihood tree of the LSU 28S rDNA of strains included in the multiple sequence alignment resembled with respect to the well supported clades (i.e. those with >80% bootstrap support) the Neighbor Joining tree in Howard et al. (2009). The isolates of *Protoceratium reticulatum* included in the phylogenetic analysis formed a monophyletic clade that further diverged into two well supported sub-clades. The two strains of *P. reticulatum* isolated in this study, A2 and H1, belong to a geographically widely distributed sub-clade that also contained the majority of the isolates (Clade A in Fig. 3 and Supplementary material Table 2). This clade contained isolates from the western and eastern North Pacific and the western and eastern North Atlantic, including the Mediterranean. One strain from South Africa represents the eastern South Atlantic and the isolates from Argentina confirm the presence of *P. reticulatum* of Clade A in the Southwest Atlantic. The second clade (Clade B in Fig. 3 and

Supplementary material Table 2) within *P. reticulatum* is represented by two isolates from the North American Atlantic coast (Florida) and one isolate from the North American Pacific coast (California). However, a wider geographic distribution of this clade may be confirmed as sampling efforts in other regions increase. The same may be the case for Clade A. The observed pattern of molecular diversification, with very low variation among sequences within sub-clades compared to the relatively high divergence among the two sub-clades, suggests that Clade A and Clade B of *P. reticulatum* correspond to two separate evolving units. The genetic distance between these two clades is about the same as that of closely related ribotypes within the *A. tamarensis/A. fundyense/A. catenella* species complex (Fig. 3). Although morphological evidence for the distinction of species in the *A. tamarensis/A. fundyense/A. catenella* complex is still not presented, it has been widely accepted that the five ribotypes represent separate species with distinct biogeographic distribution patterns and toxin properties (Scholin et al., 1994; John et al., 2003; Lilly et al., 2007; Anderson et al., 2012; Wang et al., 2014). The same might apply to the two evolutionary clades detected in *P. reticulatum*. Howard et al. (2009) considered if there are genetically distinguishable isolates of *P. reticulatum* based on YTX toxicity, but found the available data insufficient to answer this question. Indeed, in all strains from Clade A yessotoxins were found with the notable exception being one strain from New Zealand (i.e., CAWD127, Fig. 3 and Supplementary material Table 2; Rhodes et al., 2006). With regard to YTX toxicity properties of strains from Clade B, Paz et al. (2004) first found all three strains to be toxic; however, a later study by Paz et al. (2007) could not detect YTX in the same strains. However, this might eventually be explained by a loss of toxicity during the respective culture conditions (Beatriz Paz, pers. comm.¹) as a more recent study by Röder et al. (2009) again showed production of YTX in two of the same strains from Clade B previously analyzed in studies by Paz et al. (2004, 2007), i.e. CCMP404 and CCMP1720. More detailed analyses of phenotypic traits such as YTX production and mating incompatibility might show that the two ribotypic sub-clades found in *P. reticulatum* are indeed evolutionary entities with unique genetic and physiological properties – similar to PSP toxic and non-toxic species/ribotypes within the *A. tamarensis/A. fundyense/A. catenella* species complex. Detailed morphological studies might even reveal recognizable characteristics allowing for species distinction. However, already at this stage, at which the two sub-clades of *P. reticulatum* can only be identified by their ribosomal sequence divergence, future studies on the species should acknowledge the existence of these clades, e.g. when investigating YTX toxicity characteristics of the isolated strains or when studying bloom dynamics of natural populations of *P. reticulatum*. This will allow determining the specific potential of these clades for harmful algal bloom development and their role in the formation of YTX toxicity episodes.

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4.3. Spatial distribution, abundance and seasonal occurrence

The motile stage of *Protoceratium reticulatum* was quoted previously for the Southwest Atlantic by Balech (1971, 1978) in a wide area which included neritic and oceanic waters in the continental shelf of Argentina, the shelf break zone and the adjacent Atlantic basin between $\sim 37^\circ$ and 48° S. This study shows a more extensive geographic distribution ranging from Uruguayan shelf waters, where it has been also observed in samples from coastal monitoring programs (S. Méndez, pers. comm.²), to the mouth of the Magellan Strait. *P. reticulatum* has been found in coastal waters of Brazil including the southern states of Rio Grande and Santa Catarina (Cardoso, 1998; Odebrecht, 2010), and on the Pacific side its vegetative cells and cysts have been recorded in inshore waters of southern Chile (e.g. Seguel et al., 2005; Alves-de-Souza et al., 2008). We found *P. reticulatum* in a set of temperature and salinity values which include the previous regional records (Balech, 1971, 1978; Cardoso, 1998). It was recorded in neritic and estuarine environments, as is described by Steidinger and Tangen (1996), at the mouth of the Río de la Plata and off Uruguay over shelf areas of low salinity from riverine origin, and also in pelagic waters near the shelf break in a range of hydrographic conditions which comprised different water masses including the Malvinas Current (Carreto et al., 1995; Guerrero and Piola, 1997; Acha et al., 2004). Our quantitative estimations invariably showed low abundances of *P. reticulatum*. On occasions it was only present in net plankton hauls as a rare species. A low abundance generally was also observed by Balech (1971, 1978) in net plankton samples. Despite documentation as a member of planktonic communities, this species usually shows low population values (e.g. Steidinger and Williams, 1970; Seguel et al., 2005; Aasen et al., 2005), although a few records of red tides were mentioned by Reinecke (1967) and MacKenzie et al. (1998). We recorded the motile stage of *P. reticulatum* from spring to autumn, i.e. throughout the warmer period of the year, in accordance with previous records from the Southwest Atlantic (Balech, 1971, 1978; Cardoso, 1998) and the Kattegat (Hansen et al., 1996/1997). It is remarkable that during the annual study at the EPEA station the motile cells were recorded only once, in summer, meanwhile the viable resting cyst stage was present in surface sediments throughout the year. A spatio-temporal progress from north to south in the presence of the motile stage in accordance with the advance of the warm season is suggested by the distribution patterns observed from the GEF-Patagonia 1 and 2 cruises made at nearly identical positions during spring and late summer, as well as in other cruises. *P. reticulatum* cells appeared in shelf waters and the shelf break zone at the northern area in spring, while they were recorded in southern waters of the Patagonian shelf during autumn. Observations on a north–south direction in the progression of the spring bloom through analysis of seasonal variability of satellite-derived chlorophyll-*a* (Romero et al., 2006), and of paralytic shellfish toxicity maxima associated with the development and growth of *A. tamarense* (Carreto et al., 1998) were also made in this region.

Our data on viable resting cysts of *Protoceratium reticulatum* in surface sediments also showed a wide latitudinal distribution extending to the Uruguayan shelf and to the south of SJG, from its record offshore from the Río de la Plata (as *Operculodinium centrocarpum*, Wall et al., 1977) and the tidal flats of the Bahía Blanca estuary (Grill and Guersstein, 1995). In addition, living cysts were found further south ($\sim 49^\circ$ S) together with motile cells in the planktonic environment. These cysts, of apparently recent formation, suggested that an encystment process could have been occurring when we sampled. However, cell abundance was still

relatively low, a different situation than that usually observed in dinoflagellates in which encystment occurs at high cell concentration, sometimes in bloom conditions (e.g. Garcés et al., 2004), as was quoted by Reinecke (1967) for this species. The high cyst abundance and proportion in dinoflagellate assemblages found in shelf sediments from the AUCFZ and Patagonia, shows that *P. reticulatum* is one of its most important members, as has been shown in offshore pelagic and outer neritic environments of this and other regions (e.g. Wall et al., 1977; Marret and Vernal, 1997). *Protoceratium reticulatum* was mentioned as relatively abundant in plankton samples from near the EPEA station and Península Valdés (Balech, 1978). The EPEA is known as an area of development of a coastal frontal system (Carreto et al., 1995), and near Península Valdés a tidal front develops during the warm period of the year (e.g. Acha et al., 2004), and in both areas benthic cyst beds and blooms of the toxic *A. tamarense* are known to occur (Carreto et al., 1985). The high cyst values often shown by *P. reticulatum* in benthic sediments near to these frontal zones and its occasional high abundance in planktonic populations highlights the need to pay attention to the dynamics of this species in shelf areas of potential risk of shellfish contamination.

4.4. Toxins

Numerous *Protoceratium reticulatum* strains from diverse locations have been proven to produce YTX and derivatives in culture (e.g. MacKenzie et al., 1998; Ciminiello et al., 2003; Konishi et al., 2004; Paz et al., 2004, 2007, 2013; Finch et al., 2005; Krock et al., 2008a; Röder et al., 2009). To our knowledge there are only two reports of non-toxicogenic isolates of *P. reticulatum* (Rhodes et al., 2006; Paz et al., 2007), of which one (Paz et al., 2007) is contradictory to studies by Paz et al. (2004) and Röder (2010) and might be due to loss of toxicity in culture (B. Paz, pers. comm., see footnote 1). For this reason it must be taken into account that natural variability in YTX production – existing toxic and not toxic strains – have been observed in other gonyaulacoid species as well such as in *Lingulodinium polyedrum* (Reis et al., 2008) and thus may be a common trait among these species. The YTX cell quotas calculated for *P. reticulatum* from the net hauls were in the same range as the toxin cell quotas observed in the two isolates A2 and H1, and thus indicate that the factors that influence toxin production in lab conditions were similar to those in the SJG at the sampling time. The YTX cell quotas of approximately 10 pg cell^{-1} of Argentinean field samples and *P. reticulatum* isolates are in the middle range of observed *P. reticulatum* YTX cell quotas worldwide which range from 0.3 to 72 pg cell^{-1} (Paz et al., 2008). However, YTX content of field samples may also be influenced by the presence of *Gonyaulax spinifera*, which has also been shown to be a YTX producer in some areas. Whereas Rhodes et al. (2006) used only ELISA assays for the identification of YTX for a New Zealand isolate of *G. spinifera* and thus cannot attribute to its toxin profile, other authors report that *G. spinifera* YTX-profiles are dominated by homo-YTX and that yessotoxin is only a minor compound in isolates from the Adriatic Sea (Riccardi et al., 2009) and the Benguela Current (C. Chikwililwa, pers. comm.³). However, in none of the YTX containing samples, traces of homo-YTX could be detected. Furthermore, at station C43 no other known YTX-producers aside from *P. reticulatum* were present in the net haul sample. In addition there is no correlation between *G. spinifera*/*P. reticulatum* cell ratios and calculated *P. reticulatum* cell quotas. Taking all this information together, there is strong evidence that *G. spinifera* did not contribute to total YTX-toxicity in the SJG

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region. However, more research is needed to unambiguously confirm this evidence. Despite the low cell abundance of *P. reticulatum* we found, the maximum value estimated fell within the range of cell concentrations associated to varying YTX levels detected in mussels (Mallat et al., 2008; Aasen et al., 2005).

An early case of shellfish toxicity associated to this species was reported by Reinecke (1967) and Grindley and Nel (1970) in South African waters. Mouse bioassays with contaminated shellfish resulted in mouse mortalities which was suspected to be due to paralytic shellfish poisoning toxins (PSP) at that time. In contrast, mouse bioassays with plankton bloom extracts did not result in the expected mouse mortality (Grindley and Nel, 1970). Grindley and Nel used the extraction method for PSP toxins published by Sommer and Meyer (1937) which uses ethanol or methanol as extraction solvents for the toxins. However, these organic solvents would also efficiently extract lipophilic phycotoxins such as YTX from the samples. Taking into account that (1) the bloom samples were dominated by *Protoceratium reticulatum*, (2) YTX are less toxic than PSP toxins (which would explain why shellfish samples from bivalve mollusks, which can accumulate toxins to very high levels, were toxic, but not plankton extracts), and (3) organic extraction solvents were used, it is very likely that shellfish toxicities in fact were caused by YTX. Later the presence of *P. reticulatum* and its YTX-production in the same geographic area was unambiguously proven by tandem mass spectrometry (Krock et al., 2008a).

With respect to the other two known species that produce YTX, *Lingulodinium polyedrum* has been recorded in the ocean environment of the Southwest Atlantic in warm waters of the Brazil Current (Balech, 1988), but we have not found its motile cells even though their cysts were present in sediments of the EPEA station and the AUCFZ. In contrast, *Gonyaulax spinifera* is a species of widespread distribution in shelf waters of Argentina (Balech, 1988; Akselman, 1996) and we have recorded its cells at the SJG and also dinocysts of the *G. spinifera* complex as an important fraction in surface sediments. YTX were previously reported in South America from planktonic samples in north Chilean waters and in mussels from the southern coast (Yasumoto and Takizawa, 1997; Krock et al., 2009), being also detected in mussel cultures in Brazil (Schramm et al., 2010), but without knowing the causative organism. Yet in Argentina there are no information on YTX in shellfish, but our data clearly show that yessotoxins are present in plankton samples from the Patagonian area. Based on yessotoxin production and profile of local *Protoceratium reticulatum* isolates and on co-occurrence of YTX content and cell densities of this species in field samples we presented evidence that the main causal organism of YTX in the austral Southwest Atlantic is *P. reticulatum*.

5. Conclusions

Protoceratium reticulatum shows a large geographical distribution in the southern area of the Southwest Atlantic, with both its cells in the plankton habitat as well as its cysts in surface sediments. The high abundance of *P. reticulatum* viable cysts found in shelf sediments near to frontal zones such as those of Península Valdés in Patagonia and the area of the time-series station EPEA in northern Argentina, suggest the need to pay attention to the dynamics of this species in shelf areas of potential risk. In this study we have shown the presence of YTX in plankton samples of the SJG, and its production in culture by two *P. reticulatum* strains isolated from this locality. Although still there are no records of YTX in shellfish from this region, their occurrence raises the need for vigilance given the importance of commercial bivalve fisheries such as those for green-lipped mussels *Mytilus edulis* and scallops *Zygochlamys patagonica*. The ML analysis of sequences of the hypervariable region of the large subunit (LSU) 28S rDNA showed that the two clonal strains from the SJG were part of a

monophyletic clade that subdivides into two well-supported, divergent sub-clades. The sequences of the two clonal strains fell in the same clade as the majority of sequences of *P. reticulatum* and belong to a geographically wide distributed evolutionary clade. Future investigations in *P. reticulatum* need to acknowledge the cryptic (molecular) diversity in *P. reticulatum* in order to assess the specific bloom ecological and toxicity characteristics of the distinct clades.

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