

# Field observations of the dinoflagellate genus *Azadinium* and azaspiracid toxins in the south-west Atlantic Ocean

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**Abstract.** Some dinoflagellate species of the genera *Azadinium* and *Amphidoma* (Amphidomataceae) produce azaspiracids (AZA), a group of toxins responsible for gastrointestinal disorders in humans following the consumption of contaminated shellfish. In this study, we investigated the diversity, distribution and abundance of *Azadinium* and AZA from field plankton samples collected during four oceanographic expeditions that covered an extended area of the Argentine Sea during different seasons. Scanning electron microscopy analyses indicated the presence of five *Azadinium* species: *Az. dexteroporum*, *Az. luciferelloides*, *Az. obesum*, *Az. asperum* and *Az. cf. poporum*. *Azadinium*-like cells were frequently found and were even an abundant component of plankton assemblages, showing a wide latitudinal distribution, from ~38 to ~55.5°S, and occurring in a wide temperature and salinity range. High cell densities (up to 154 000 cells L<sup>-1</sup>) occurred in northern slope and external shelf waters during spring. AZA-2 was detected in net samples from the 20- to 200-µm fractions by tandem mass spectrometry–liquid chromatography analysis, suggesting a transfer of AZA through the food web. Our results contribute to the knowledge of the worldwide occurrence of *Azadinium* species and AZA, and highlight the importance of amphidomatacean species as a potential source of AZA shellfish poisoning in the south-west Atlantic Ocean.

**Additional keywords:** *Azadinium asperum*, *Azadinium dexteroporum*, *Azadinium luciferelloides*, *Azadinium obesum*, *Azadinium cf. poporum*, azaspiracid-2.

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## Introduction

The consequences of harmful algal blooms related to phyco-toxin transfer through the food web, which includes effects on public health, fauna mortality and negative repercussions on the marine ecosystem, seem to have increased since the 1960s. Similarly, new phycotoxins and toxin-producing species have been discovered in recent years; of these, the most recent examples are the dinoflagellate family Amphidomataceae (which includes species of *Azadinium* and *Amphidoma*) and their toxins, azaspiracids (AZAs; Lassus *et al.* 2016). To date, 26 AZAs have been reported from dinoflagellate origin (Krock *et al.* 2019). Some of these have been known since the 1990s as the source of human intoxications by AZAs, a syndrome called azaspiracid shellfish poisoning (Satake *et al.* 1998) that includes severe gastrointestinal disorders after ingestion of contaminated seafood (Twiner *et al.* 2014). At present, the genus *Azadinium* consists of 13 species (Tillmann *et al.* 2014a; Tillmann and Akselman 2016; Luo *et al.* 2017; Tillmann 2018), of which 11

have been tested for AZA production and three (*Azadinium spinosum*, *Azadinium dexteroporum* and *Azadinium poporum*) have shown the capacity to synthesise AZA (Tillmann *et al.* 2009; Krock *et al.* 2012; Rossi *et al.* 2017). Only two of the 16 species of the genus *Amphidoma* have been tested for the production of AZA, with *Amphidoma languida* (Krock *et al.* 2012), but not *Amphidoma parvula* (Tillmann *et al.* 2018a), proven to produce AZA.

Most *Azadinium* species are small (<20 µm) and their morphological identification requires detailed scanning electron microscopy (SEM) to reveal plate patterns and other diagnostic characteristics at the species level, such as the presence or absence of an antapical spine, the location of the ventral pore and some specific characteristics of diagnostic plates, such as Plate 2a (Tillmann *et al.* 2009, 2010, 2012a, 2012b, 2014a). Consequently, identification of *Azadinium* is complex and thus little is known regarding their worldwide distribution. However, the wide distribution of AZA analogues

detected in shellfish samples (e.g. Braña Magdalena *et al.* 2003; Taleb *et al.* 2006; Elgarch *et al.* 2008; Vale *et al.* 2008; Álvarez *et al.* 2010; Yao *et al.* 2010; Trainer *et al.* 2013; Krock *et al.* 2014; Bacchiocchi *et al.* 2015; Turner and Goya 2015) indicates that AZA-producing Amphidomataceae species are cosmopolitan.

Records of Amphidomataceae for the southern Atlantic continental shelf may be traced back to the mid-1970s when Campodónico and Guzmán (1974) described a dense bloom ( $6.8 \times 10^5$  cells  $L^{-1}$ ) of *Amphidoma* sp. in the Magallanes Strait that, interestingly, coincided with a mild case of human intoxication after shell consumption. However, Campodónico and Guzmán (1974) did not provide micrographs, drawings or any morphological description other than size (and that cells have spines) in their report, so the identification of the causative species as *Amphidoma* is rather doubtful and needs confirmation. In the Argentine Sea, the first record of Amphidomataceae corresponds to one cell of an unidentified species of *Amphidoma* found by Balech (1988) in northern slope waters ( $\sim 39^\circ S$ ,  $52^\circ W$ ). After that, two Amphidomataceae spring blooms were registered in 1990 and 1991 from slope waters adjacent to Mar del Plata, Buenos Aires Province ( $\sim 39^\circ S$ ) and from the middle shelf of an area neighbouring El Rincón, Buenos Aires Province ( $\sim 40^\circ S$ ; Akselman and Negri 2012). The 1990 bloom generated water discolouration that lasted for 1 week and covered an area of  $\sim 60$  km<sup>2</sup>. The maximum cell density of  $9 \times 10^6$  cells  $L^{-1}$  (Akselman and Negri 2012) represents the biggest *Azadinium* bloom thus far. The presence of an antapical spine and a pyrenoid in the cells led Akselman and Negri (2012) to identify the cells as *Azadinium* cf. *spinosum*, one of the three *Azadinium* species described at that time. However, some taxonomic characteristics, such as the location of the ventral pore, could not be observed. Later, a third bloom was recorded in late winter 1998 in the same area as the 1990 bloom, again causing water discolouration; the cells in this case were also characterised as *Az. cf. spinosum* (Akselman *et al.* 2014). However, a re-examination of the 1990 bloom samples with SEM revealed that a newly described species, *Azadinium luciferelloides*, was the dominant species in the community (Tillmann and Akselman 2016), and one additional species of *Azadinium* (*Azadinium asperum*) and three new *Amphidoma* species (*Am. trioculata*, *Am. cyclops* and *Am. alata*), in addition to the SEM record of *Am. languida* were also described (Tillmann 2018). The first *Azadinium* strains from Argentina were *Az. poporum*, established from sediment samples collected from a highly productive area of coastal waters off Buenos Aires Province (El Rincón), with toxin analysis showing profiles consisting exclusively of AZA-2 (Tillmann *et al.* 2016). Based on a survey in 2015 across El Rincón to the outer shelf area, light microscopy and polymerase chain reaction (PCR) analysis revealed that Amphidomataceae were widely present and abundant in spring plankton communities, but AZA could not be detected in the plankton (Tillmann *et al.* 2019). From this survey, several strains of *Azadinium dalianense* and *Az. spinosum* and a strain of the newly described *Am. parvula* were established. Toxin analyses showed that all *Az. dalianense* and most *Az. spinosum* strains were non-toxicogenic, except for one strain that produced only AZA-2 (Tillmann *et al.* 2019). Regarding detection of AZAs in Argentinean mussels, low levels of AZA-2 were

detected in shellfish from Santa Teresita and San Clemente del Tuyú, Buenos Aires Province (Turner and Goya 2015).

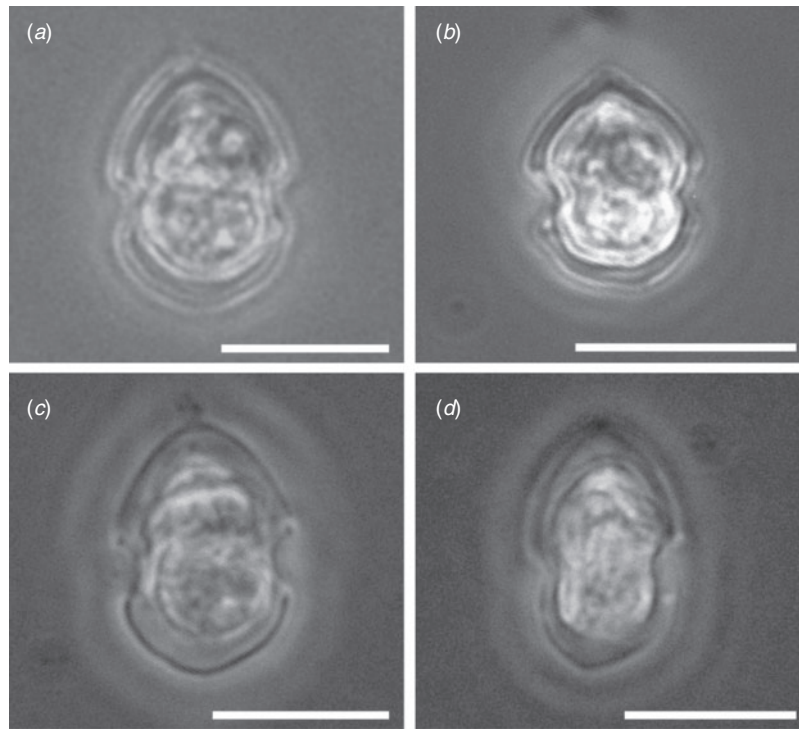
Despite these previous reports of *Azadinium* and AZAs, information about their diversity and distribution remains limited and restricted to the northern Argentine Sea. Even though the identification and quantification of toxicogenic Amphidomataceae species is challenging, particularly because both toxicogenic and non-AZA-producing species occur in the same area (Tillmann *et al.* 2010, 2011, 2012b, 2019), it is important to determine the occurrence of *Azadinium* and *Amphidoma* species and the presence of AZA in the south-west Atlantic. The aim of the present study was to analyse the abundance and spatiotemporal distribution of Amphidomataceae species in field populations from an extended area of the Argentine Sea coupled with field sample AZA analysis and electron microscopy for species identification.

## Material and methods

### Field sampling protocols

The continental shelf waters of the Argentine Sea were sampled during four oceanographic expeditions. Expedition 1 (E1) was conducted in austral autumn onboard *R/V Puerto Deseado* from 30 March to 14 April 2012. In all, 46 stations were sampled between  $\sim 38$  and  $56^\circ S$ . The second expedition (E2) was conducted in late austral summer on *R/V Bernardo Houssay* from 11 to 22 March 2013, with 24 sampling stations located between  $\sim 39$  and  $43^\circ S$ . This cruise was divided in two legs, K1 and K2, which comprised 8 and 16 sampling stations respectively. The third expedition (E3) was conducted in austral spring aboard *R/V Puerto Deseado* from 26 October to 9 November 2013, with 47 sampling stations located between  $\sim 40$  and  $47^\circ S$ . Finally, the fourth expedition (E4) was conducted on *R/V Bernardo Houssay* in austral summer from 6 to 12 January 2016, with seven sampling stations located between  $\sim 42$  and  $55^\circ S$ . The conductivity, temperature and depth (CTD) data were available throughout all expeditions, except from K2 of E2, during which no CTD measurements were made. During this leg, only surface water temperature was measured with a multiparameter probe (Model WQC; TOA-DKK Corporation, Tokyo, Japan).

Plankton samples were collected by vertical net tows through the upper 20 m of the water column using a 20- $\mu m$  mesh Nitex net 60 cm in diameter for both taxonomic and phycotoxin analysis. Each net haul concentrate was adjusted to 1 L with 0.2  $\mu m$  of filtered seawater. A 100-mL aliquot was fixed with acidic Lugol's iodine solution for species identification and enumeration. The rest was sequentially filtered through Nitex mesh of 200, 50 and 20  $\mu m$  in polyvinyl chloride (PVC) cylinders by gravity filtration. The particulate material retained on each mesh was resuspended in 40 mL of filtered seawater and transferred to 50-mL centrifuge tubes. Each size fraction concentrate was split into two 20-mL aliquots for lipo- and hydrophilic toxin analyses. The aliquots were centrifuged at 3220g for 15 min at room temperature. The pellets were stored at  $-20^\circ C$  for later toxin analysis. During all four expeditions, Niskin bottle samples were taken from depths of 3 and 10 m, mixed in equal volume and fixed with acidic Lugol's iodine solution for quantitative analysis of phytoplankton. For AZA analysis, 2- or 3-L samples of this mixture were filtered through a 3- $\mu m$



**Fig. 1.** Light microscopy images of *Azadinium*-like cells found in Lugol-fixed bottle samples. Scale bars: 10  $\mu\text{m}$ .

(E1–E3) or 5- $\mu\text{m}$  (E4) pore polycarbonate filter. Filters were placed with their back to the inner wall of a 50-mL centrifuge tube (Sarstedt, Nümbrecht, Germany) and kept at  $-20^{\circ}\text{C}$  for less than 2 months until analysis.

#### Plankton analysis

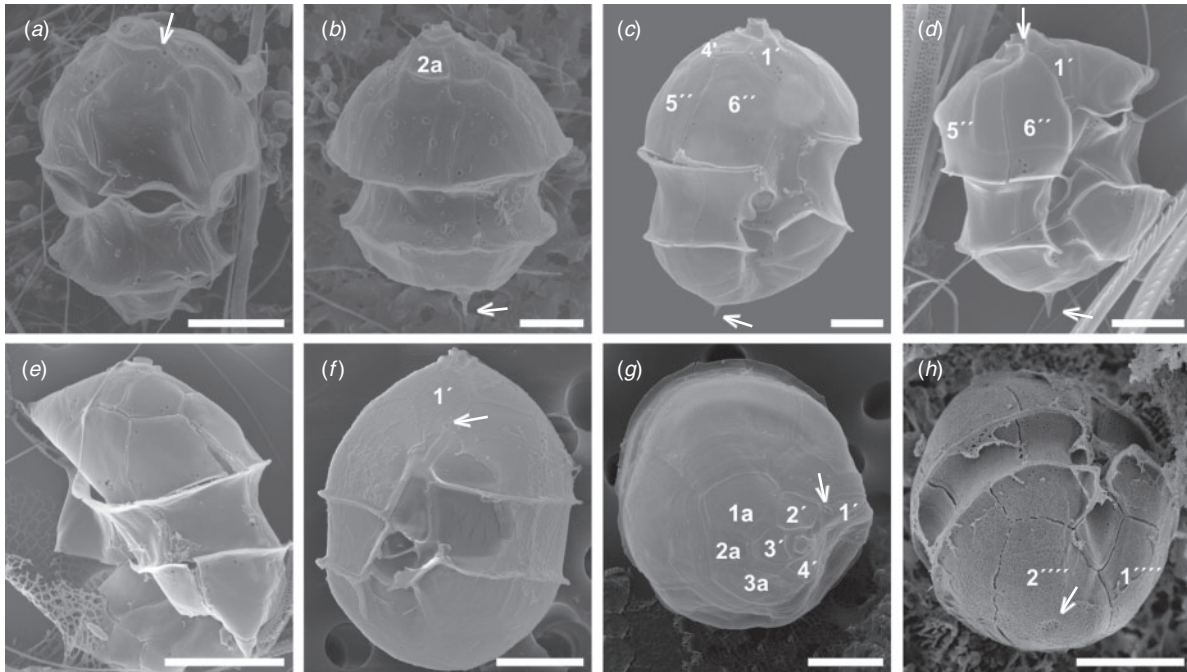
The abundance of nano- ( $\sim 5$ – $20\ \mu\text{m}$ ) and microplankton ( $20$ – $200\ \mu\text{m}$ ) was estimated using the Utermöhl (1958) inverted microscope method. Subsamples (50 mL) from the mixed water obtained using Niskin bottles were left to settle for 24 h in a composite sedimentation chamber before cell counting. At least 300 cells of the dominant taxa were counted in random fields or in half the chamber to estimate general plankton composition, whereas the whole chamber bottom was scanned to count less abundant species. Plankton taxa from the 5- to  $200\text{-}\mu\text{m}$  size range, including photosynthetic and heterotrophic protists, were identified to the highest possible taxonomic level (i.e. species). However, some small flagellates ( $\sim 5$ – $10\ \mu\text{m}$ ) that lost their flagella during fixation were included in a single group as ‘small unidentified flagellates’. In particular, dinoflagellate abundance includes cells with and without plastids. During plankton counting, small thecate dinophytes that were between  $\sim 7$  and  $18\ \mu\text{m}$  long and had a pointed apex and a wide cingulum (Fig. 1) were counted in the category of *Azadinium*-like cells, but it has to be kept in mind that such a category probably also includes *Amphidoma* sp. and may also contain small *Heterocapsa* species; thus, cell abundances may be overestimated. The limit of detection of this method was  $20\ \text{cells L}^{-1}$  for the three expeditions.

Further morphological examination of selected samples was conducted with three SEMs: Jeol JSM-6360 LV SEM (JEOL, Tokyo, Japan), Carl Zeiss NTS SUPRA 40 (Zeiss, Oberkochen, Germany) and FEI Quanta FEG 200 (FEI, Eindhoven, Netherlands). Aliquots (50 mL) of bottle samples were filtered through  $0.2\text{-}\mu\text{m}$  polyamide filters for SEM analyses. The material on the filters was dehydrated by serial ethanol treatment (25, 50, 75 and 100%) and final critical point dehydration (BAL-TEC CPD-30; BAL-TEC, Balzers, Liechtenstein). Specimens were sputter-coated with Au with an iron sputter fine coat Jeol JFC 1.100 (JEOL) before examination. The following samples were analysed by SEM: I9, I13 and I15 from E1; 3K1, 9K2, 10K2 and 20K2 from E2; 1, 3–7, 25, 43 and 44 from E3; and 4 and 5 from E4.

#### Toxin analysis

Cell pellets obtained from net samples were suspended in  $500\ \mu\text{L}$  methanol, and subsequently homogenised with  $0.9\ \text{g}$  of Lysing Matrix D (Thermo Savant, Illkirch, France) by reciprocal shaking at maximum speed ( $6.5\ \text{m s}^{-1}$ ) for 45 s in a Bio101 FastPrep instrument (Thermo Savant, Illkirch, France). After homogenisation, samples were centrifuged at  $16\ 100g$  for 15 min at  $4^{\circ}\text{C}$ . The supernatant was transferred to a spin filter (pore size  $0.45\ \mu\text{m}$ ; Millipore Ultrafree, Eschborn, Germany) and centrifuged at  $800g$  for 30 s at room temperature, followed by transfer to autosampler vials. Toxin concentrations are expressed as nanograms per net tow.

For bottle sample analysis of AZA, polycarbonate filters were repeatedly rinsed with  $500$ – $1000\ \mu\text{L}$  of methanol until



**Fig. 2.** Scanning electron microscopy images of *Azadinium* cells from the Argentine Sea. (a, b) *Azadinium dexteroporum*. (a) Lateral view; note the location of the ventral pore (vp) at the right margin of the pore plate (arrow). (b) Dorsal view; note the presence of an antapical spine (arrow) and the concave shape of Plate 2a. (c–e) *Azadinium luciferelloides*. (c, d) Ventral views; note the presence of an antapical spine (arrows) and the location of the vp at the right side of the pore plate (d; arrow). (e) Dorsal view. (f) *Azadinium obesum* ventral view; note the position of the vp at the left side of Plate 1' (arrow). (g) *Azadinium asperum* apical view; note the location of the vp in the left region of Plate 1' at the end of the suture between Plates 1' and 2' (arrow). (h) *Azadinium cf. poporum* antapical view; note the group of pores in Plate 2'''' (arrow). Scale bars: 2  $\mu\text{m}$  (a–d); 5  $\mu\text{m}$  (e–h).

complete discolouration of the filters. The methanolic extracts were transferred to a spin filter (pore size 0.45  $\mu\text{m}$ ; Millipore Ultrafree), and centrifuged at 800g for 30 s at 4°C, followed by transfer to autosampler vials. AZA concentrations in samples were determined using liquid chromatography–tandem mass spectrometry (LC-MS/MS), as described by Luo *et al.* (2016).

The following AZA analogues were included in bottle and net samples analysis: 1, 2, 11, 33–43, 54–59 and 62.

## Results

### Species identified

Five *Azadinium* species were identified by SEM analysis of field samples: *Az. dexteroporum* ( $n = 15$ ), *Az. luciferelloides* ( $n = 5$ ), *Az. obesum* ( $n = 4$ ), *Az. asperum* ( $n = 1$ ) and *Az. cf. poporum* ( $n = 3$ ). No *Amphidoma* species were identified by SEM analysis of field samples.

The identification of *Az. dexteroporum* was based on the location of the ventral pore (vp) at the end of an asymmetrical elongation of the pore plate (Po) and by the particular shape of the intercalary Plate 2a, which was concave, forming a depression on the theca (Fig. 2a, b). *Az. luciferelloides* was identified by the position of the vp at the right side of the Po and by the antapical spine. In addition, the presence or absence and arrangement of thecal pores (diameter 0.05–0.15  $\mu\text{m}$  (mean 0.1  $\mu\text{m}$ );  $n = 27$ ) was also consistent with the species description (Fig. 2c–e). The identification of *Az. obesum* was based on the location of the vp at the posterior left side of Plate 1', by the

absence of an antapical spine and by a slightly larger size of the cells (Fig. 2f). *Az. asperum* was characterised by the presence of an antapical spine and by the location of the vp in the anterior left region of Plate 1'. *Az. asperum* could also be differentiated from other *Azadinium* by the presence of a rugged theca (Fig. 2g). Finally, a few cells without an antapical spine and with a group of 5–10 pores on Plate 2'''' were identified as *Az. cf. poporum* (Fig. 2h).

### *Azadinium* abundance and distribution

#### Autumn expedition (E1)

*Azadinium*-like cells were found in 15 of 47 bottle samples at cell densities ranging between 20 and 4000 cells  $\text{L}^{-1}$  and representing between 0.4 and 31 % of total dinoflagellate abundance (mean 10%). Maximum abundance of *Azadinium*-like cells was found in the southern sampling area (55°S), in shelf waters from Tierra del Fuego Province and Isla de los Estados (Fig. 3). At these stations (I9, I13 and I15), concentrations of total planktonic protists reached  $1.1\text{--}5.6 \times 10^5$  cells  $\text{L}^{-1}$ , and these were primarily represented (i.e. 69–97%) by small (~5–10  $\mu\text{m}$ ) unidentified flagellates. Dinoflagellate abundance in those samples was  $9\text{--}14 \times 10^3$  cells  $\text{L}^{-1}$  (which represented 2–11% of total planktonic protists) and dinoflagellates were primarily represented by unidentified small gymnodinioids ( $2.5\text{--}8.1 \times 10^3$  cells  $\text{L}^{-1}$ ). *Azadinium*-like cells were present at lower densities ( $20\text{--}10^3$  cells  $\text{L}^{-1}$ ) in another seven samples from that area, which showed a similar plankton composition as reported for

Stations I9, I13 and I15. However, in five samples of the San Jorge Gulf, where *Azadinium*-like cells were also found, plankton composition was considerably different from that of the southern stations. In four of those samples (C43, C43N, C44 and C45) dinoflagellates were the dominant group, being present at densities between  $6 \times 10^3$  and  $34 \times 10^3$  cells  $L^{-1}$ , and primarily represented by the genus *Tripos*. At Station P45B, diatoms dominated the community, with an abundance of  $6 \times 10^3$  cells  $L^{-1}$  and primarily represented by *Thalassiosira* sp.

During the whole cruise, *Azadinium*-like cells were detected over a temperature range of 6–14°C and at salinities ranging from 31.2 to 34.2. Maximum *Azadinium*-like cell densities were found at temperatures between 7.8 and 8.6°C and at salinities of 32.6–33.7. The temperature range for this expedition was 6–19°C and the salinity range was 31.2–34.2.

Through the concentration of bottle samples from Stations I9, I13 and I15, obtained in the southern Argentine Sea (~55°S), it was possible to identify two *Azadinium* species: *Az. dexteroporum* and *Az. cf. poporum* (Fig. 3).

#### Late summer expedition (E2)

*Azadinium*-like cells were detected in 22 of 25 bottle samples, with cell abundances ranging from 20 to 19 000 cells  $L^{-1}$  and representing 0.3–53% of total dinoflagellate abundance (mean 17%). The maximum abundance of *Azadinium*-like cells was found in southern San Matías Gulf, at Station 10K2 (Fig. 4). At this station, total phytoplankton abundance was  $3.4 \times 10^5$  cells  $L^{-1}$ , and there was a high contribution of dinoflagellates (34%), primarily represented by *Prorocentrum cordatum*. Unidentified small (<10 µm) flagellates were dominant at all samples with *Azadinium*-like cells, except for samples corresponding to Valdés Peninsula, where cryptophytes were the dominant group. In addition, although abundance was lower (<5000 cells  $L^{-1}$ ), *Azadinium*-like cells represented more than 30% of total dinoflagellates at Stations 6K1 (southern Buenos Aires Province), 20K2, 21K2 (San Matías Gulf) and 28K2 (Valdés Peninsula), with total planktonic protist abundances of  $0.7$ – $1.8 \times 10^5$  cells  $L^{-1}$  and dinoflagellate densities of  $0.02$ – $0.1 \times 10^5$  cells  $L^{-1}$ . Diatoms, primarily represented by *Guinardia delicatula*, dominated at Station 20K2.

*Azadinium*-like cells were found in a temperature range of 15–19°C and at salinities ranging from 31.9 to 34.1. Maximum *Azadinium*-like cell density was found at 16.2°C. The temperature range during E2 was 14–19°C and salinity ranged between 31.9 and 34.1.

It was not possible to identify *Azadinium* cells from this expedition to species level because SEM analyses did not enable enough taxonomic features to be seen to assign cells to a particular species.

#### Spring expedition (E3)

*Azadinium*-like cells were found in 26 of 44 bottle samples, at cell densities ranging from 20 to  $1.5 \times 10^5$  cells  $L^{-1}$  and representing between 0.05 and 87% of total dinoflagellate abundance (mean 21%). In 10 samples, *Azadinium*-like cells accounted for more than 30% of total dinoflagellate abundance. Eight of those samples were from stations located in slope waters (Stations 1, 4, 5, 6, 7, 39, 43 and 44) with total

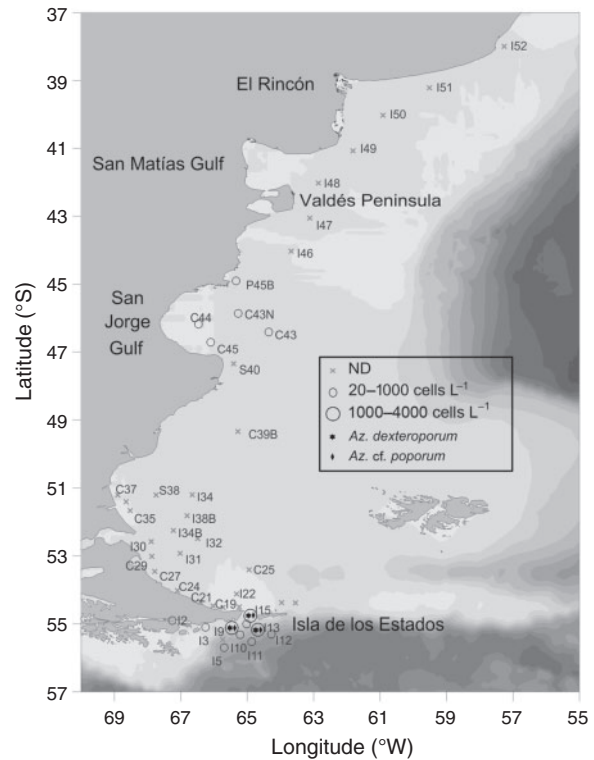


Fig. 3. Distribution and abundance of the *Azadinium*-like cells in bottle samples from the autumn 2012 expedition (Expedition 1). ND, not detected.

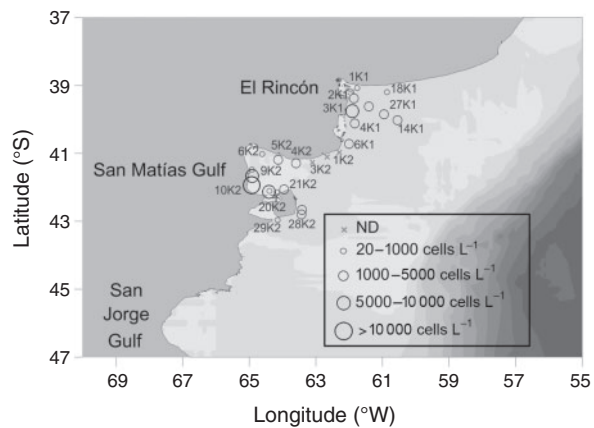
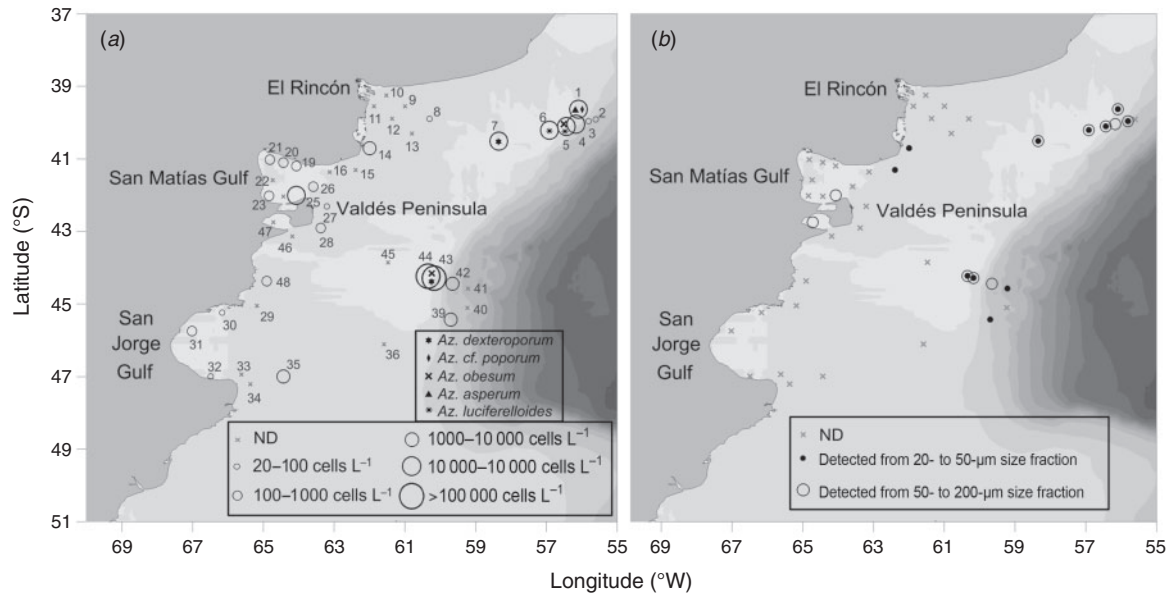


Fig. 4. Distribution and abundance of *Azadinium*-like cells in bottle samples from the late summer 2013 expedition (Expedition 2). ND, not detected.

plankton abundances of  $0.8$ – $40.4 \times 10^5$  cells  $L^{-1}$  and dominated by diatoms (primarily *Hemiaulus* sp., *Helicotheca tamensis* and *Thalassiosira* sp.), with the exception of samples from Station 6, which was dominated by dinoflagellates (primarily *Protoperidinium* sp., with an abundance of  $0.2 \times 10^5$  cells  $L^{-1}$ ), and Station 39, which was dominated by small unidentified flagellates. At the other two stations, Stations 14 and 25, total planktonic protists abundance was  $120.6 \times 10^5$  and



**Fig. 5.** (a) Distribution and abundance of *Azadinium*-like cells in bottle samples and (b) azaspiracid (AZA)-2 distribution in net samples from the spring 2013 expedition (Expedition 3). ND, not detected.

$34.7 \times 10^5$  cells  $L^{-1}$  respectively, and the dominant groups were small unidentified phytoflagellates and haptophytes respectively. Maximum abundance of *Azadinium*-like cells was detected at Stations 43 and 44 located in slope waters between 40 and 44°S (Fig. 5a), in which the maximum abundance of *Gyrodinium fusus* ( $10\text{--}11 \times 10^3$  cells  $L^{-1}$ ) was also recorded. At Stations 14 and 35, moderate densities ( $1\text{--}10 \times 10^3$  cells  $L^{-1}$ ) of *Azadinium*-like cells were found and total plankton abundance was  $120.7 \times 10^5$  and  $10.5 \times 10^5$  cells  $L^{-1}$  respectively. In samples from both Stations 14 and 35, unidentified small flagellates were dominant ( $120.0 \times 10^5$  and  $9.2 \times 10^5$  cells  $L^{-1}$  respectively), but a high abundance ( $37.5 \times 10^3$  cells  $L^{-1}$ ) of the diatom *Asterionellopsis* sp. was found at Station 14 and prasinophytes, with a cell density of  $100 \times 10^3$  cells  $L^{-1}$ , were the second-most abundant group at Station 35.

*Azadinium*-like cells were present at temperatures ranging from 8 to 13.5°C and at salinities ranging from 33 to 34.1. Maximum *Azadinium*-like cell densities were detected between 9.4 and 9.7°C and at a salinity of 33.7. The temperature range for this expedition was 7–15°C and the salinity range was 32.9–4.2.

SEM analyses enable identification of five species: *Az. obesum*, *Az. dexteroporum*, *Az. luciferelloides*, *Az. asperum* and *Az. cf. poporum*. The presence of *Az. asperum* and *Az. cf. poporum* was confirmed at Station 1 and *Az. luciferelloides* was confirmed at Stations 5 and 6, all corresponding to northern slope waters (~40°S). *Az. obesum* was found at Stations 5 and 43, whereas *Az. dexteroporum* was detected at Stations 7, 43 and 44, all from northern (~40°S) and central (~44°S) slope waters (Fig. 5a).

*Azadinium*-like cells were also found by light microscopy in three net samples from the 20- to 50- $\mu$ m size fractions of Stations 1, 6 and 26, and in one sample of the 50- to 200- $\mu$ m fraction of Station 1.

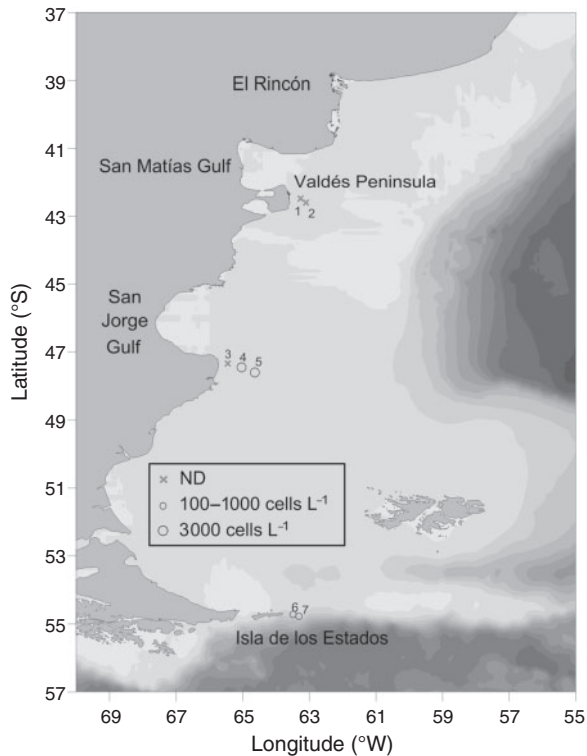
#### Early summer expedition (E4)

*Azadinium*-like cells were found in four of seven bottle samples, with densities between 100 and 3000 cells  $L^{-1}$  and representing 1.5–10% of total dinoflagellate abundance (mean 4%). Maximum cell densities were found in southern San Jorge Gulf (47°S; Fig. 6) in samples with a total phytoplankton abundance of  $10.8\text{--}17.5 \times 10^5$  cell  $L^{-1}$  and dominated by small unidentified flagellates. At those stations dinoflagellates reached high abundances ( $34\text{--}128 \times 10^3$  cells  $L^{-1}$ ), primarily due to the heterotrophic *G. fusus*. At the other two stations with *Azadinium*-like cells, Stations 6 and 7, small unidentified flagellates (<10  $\mu$ m) and diatoms (primarily represented by *Pseudo-nitzschia* species) were dominant respectively.

*Azadinium*-like cells were found at temperatures ranging from 5 to 11.5°C and at salinities ranging from 32.9 to 34.1. Maximum *Azadinium*-like cell densities were found at temperatures between 11.1 and 11.4°C and at a salinity of 33. The temperature range of the whole expedition was 5–15°C and salinity ranged between 32.9 and 34.0. The low cell abundances did not allow further SEM analysis for species identification.

#### Azaspiracids

For all four expeditions, AZA analysis of bottle samples were negative with the limits of detection of 43, 37–444, 59–89 and 26–132  $\mu$ g  $L^{-1}$  for E1, E2, E3 and E4 respectively. However, AZA-2 was detected in spring 2013 (E3) in plankton pellets obtained from 11 net samples corresponding to the 20- to 50- $\mu$ m size fraction and from another 11 net samples corresponding to the 50- to 200- $\mu$ m size fraction. AZA-2 was detected at stations located in slope waters from 39 to 40°S and from 44 to 45°S, from San Matías and Nuevo Gulfs and southern Buenos Aires Province (Fig. 5b). AZA-2 abundances ranged between 29 and 1244  $\mu$ g per net tow. Plankton net sample AZA concentrations



**Fig. 6.** Distribution and abundance of *Azadinium*-like cells in bottle samples from the early summer 2014 expedition (Expedition 4). ND, not detected.

from the other three expeditions were below the limit of detection of  $\sim 30$  pg per net tow.

## Discussion

### *Azadinium* diversity

The microscopic identification of *Azadinium* species in field samples is difficult, especially when cell densities are low and different species occur in the same area. Moreover, species identification in most cases is impossible using light microscopy only (Tillmann *et al.* 2010, 2011, 2012b). In the present study it was possible to identify five *Azadinium* species by concentration of Niskin bottle samples and further SEM analysis.

*Az. obesum* was described based on a North Sea strain (Tillmann *et al.* 2010). This species is known to occur in the North Atlantic (Tillmann *et al.* 2018b) and north-east Pacific (Kim *et al.* 2017), and the present record is the first detection of *Az. obesum* in the Southern Hemisphere. None of the cultured strains of this species produced AZA analogues detectable by LC-MS/MS (Tillmann *et al.* 2010, 2018b; Kim *et al.* 2017).

*Az. luciferelloides* was recently described based on a field sample collected during an Amphidomataceae bloom that occurred in the 1990s in Argentinean slope waters. This species was the most abundant of all Amphidomataceae cells in that sample (Tillmann and Akselman 2016). Although there are another three *Azadinium* species (*Az. caudatum*, *Az. dexteroporum* and *Az. concinnum*) that, like *Az. luciferelloides*, have a vp in the right region of the Po (Tillmann and Akselman 2016),

*Az. caudatum* is significantly larger than *Az. luciferelloides* and has a different cell shape, *Az. dexteroporum* has a vp located at the end of an asymmetrical elongation of the pore plate and *Az. concinnum* has very large and symmetric precingular plates and very small apical plates (Tillmann *et al.* 2014a). The size range of *Az. luciferelloides* estimated here (7.4–13.6  $\mu\text{m}$  long, 4.8–10.2  $\mu\text{m}$  wide) is similar to the size reported in the species description (6.4–14.1  $\mu\text{m}$  long, 7.6–11.6  $\mu\text{m}$  wide; Tillmann and Akselman 2016). *Az. luciferelloides* was detected during the spring expedition in southern slope waters at 40°S in two samples in which it was the only species identified and the abundance of *Azadinium*-like cells was high ( $30 \times 10^3$  cells  $\text{L}^{-1}$ ). This confirms that this species is an important member of the *Azadinium* spring communities in Argentina. Interestingly, *Az. luciferelloides* has a wide distribution, ranging from cool slope Argentinean waters (Tillmann and Akselman 2016; present study) to tropical Brazilian waters (Cavalcante *et al.* 2018). No cultured strain of the species is available yet, so its potential for AZA production is unclear.

During E1 and E3, the presence of *Az. dexteroporum* was confirmed for slope waters close to the site of the 1991 bloom ( $\sim 38^\circ\text{S}$ ) and in the southern Argentine Sea (44 and 55°S). The species *Az. dexteroporum* was described based on a strain isolated from the Gulf of Naples in the Mediterranean Sea (Percopo *et al.* 2013). Tillmann and Akselman (2016) found this species as part of the 1991 bloom in the Argentine Sea, although it was present in low abundance relative to total Amphidomataceae cells. In accordance with Tillmann and Akselman (2016), the cells observed in the present study had a vp located closer to the Po than specimens from the Mediterranean Sea (Percopo *et al.* 2013). The toxin-producing potential of *Az. dexteroporum* remains uncertain at the moment: culture of the Mediterranean type showed production of AZA-35 in addition to six compounds that represented new additions to the AZA group of toxins (Rossi *et al.* 2017), whereas no AZAs were detected in an Iceland strain isolated from the Irminger Sea (Tillmann *et al.* 2015). None of the recently described AZA analogues (i.e. AZA-54–58) produced by Mediterranean isolates of *A. dexteroporum* (Rossi *et al.* 2017) were found in the present study, despite the presence of this species in field samples. Therefore, the Argentinean populations of *A. dexteroporum* do not appear to be AZA-producing strains, similar to the strains isolated from Iceland (Tillmann *et al.* 2015), but confirmation by testing local strains is needed.

The latest described *Azadinium* species, *Az. asperum* (Tillmann 2018), was detected in the present study in slope waters of the northern sampling area in accordance with its type locality (Tillmann 2018). No cultured strain of the species is available yet, so the potential for producing AZA is unclear.

In SEM preparations of field samples, most amphidomatacean cells were impossible to identify at a species level because the plates are wrinkled, cells are deformed or oriented in such a way on the filter that important morphological details are hidden. Among those cells, three specimens visible in antapical view attracted our attention because of the presence of a distinct field of pores and a lack of an antapical spine on the second antapical plate. Such a group of pores was also found in Argentinean *Az. poporum* strains (Tillmann *et al.* 2016) and in *Az. poporum* from China (Gu *et al.* 2013), the Gulf of Mexico

(Luo *et al.* 2016) and the Mediterranean (Luo *et al.* 2017). Although no other diagnostic traits for *Az. poporum* (i.e. position of the vp, presence of four apical plates to differentiate *Az. poporum* from *Az. dalianense*) were visible for any of the Argentinean field sample specimens, we used the presence of a group of pores and the absence of an antapical spine as evidence to classify these specimens as *Az. cf. poporum*. Such cells were detected in samples from external shelf waters at 55 and 40°S during E1 and E3 respectively. The presence of *Az. poporum* in Argentina is known from local strains (Tillmann *et al.* 2016) and from positive PCR assays using specific primers and probes (Tillmann *et al.* 2019).

Information from the present study together with previous reports (Tillmann and Akselman 2016; Tillmann *et al.* 2016, 2019; Tillmann 2018) highlights the Argentine Sea as an area harbouring quite a diverse range of *Azadinium* species, considering that 6 of the 13 species are recorded from this region and that 2 were actually described from Argentinean field samples. Conversely, other Amphidomataceae species (i.e. *Am. languida*, *Am. parvula*, *Am. cyclops*, *Am. alata*, *Am. trioculata*, *Az. spinosum* and *Az. dalianense*) previously recorded from the exceptional 1990–91 blooms or by the establishment of cell cultures and the use of molecular markers (Tillmann and Akselman 2016; Tillmann *et al.* 2018a, 2019; Tillmann 2018) were not detected in the present study. The fact that they were not found in SEM analysis of our field samples could be the result of their presence in very low abundances.

#### *Azadinium* and AZA distribution

Despite the importance of detecting and quantifying toxigenic species in the field, studies about the spatial and temporal dynamics of *Azadinium* species are scarce. In the Argentine Sea, there are few Amphidomataceae reports since the work of Akselman and Negri (2012) and Akselman *et al.* (2014), who registered three intense *Azadinium* blooms in the 1990s, two of them during spring and one during late winter, in middle shelf waters adjacent to the front of El Rincón and in slope waters. Recently, a spring expedition revealed Amphidomataceae densities up to  $2.8 \times 10^5$  cells L<sup>-1</sup> at El Rincón in 2015 (Tillmann *et al.* 2019). The idea that the shelf front of the Argentine coast in spring is a hot spot for amphidomatacean species is supported by the present results, with maximum *Azadinium*-like cell densities found in northern slope and external shelf waters during spring.

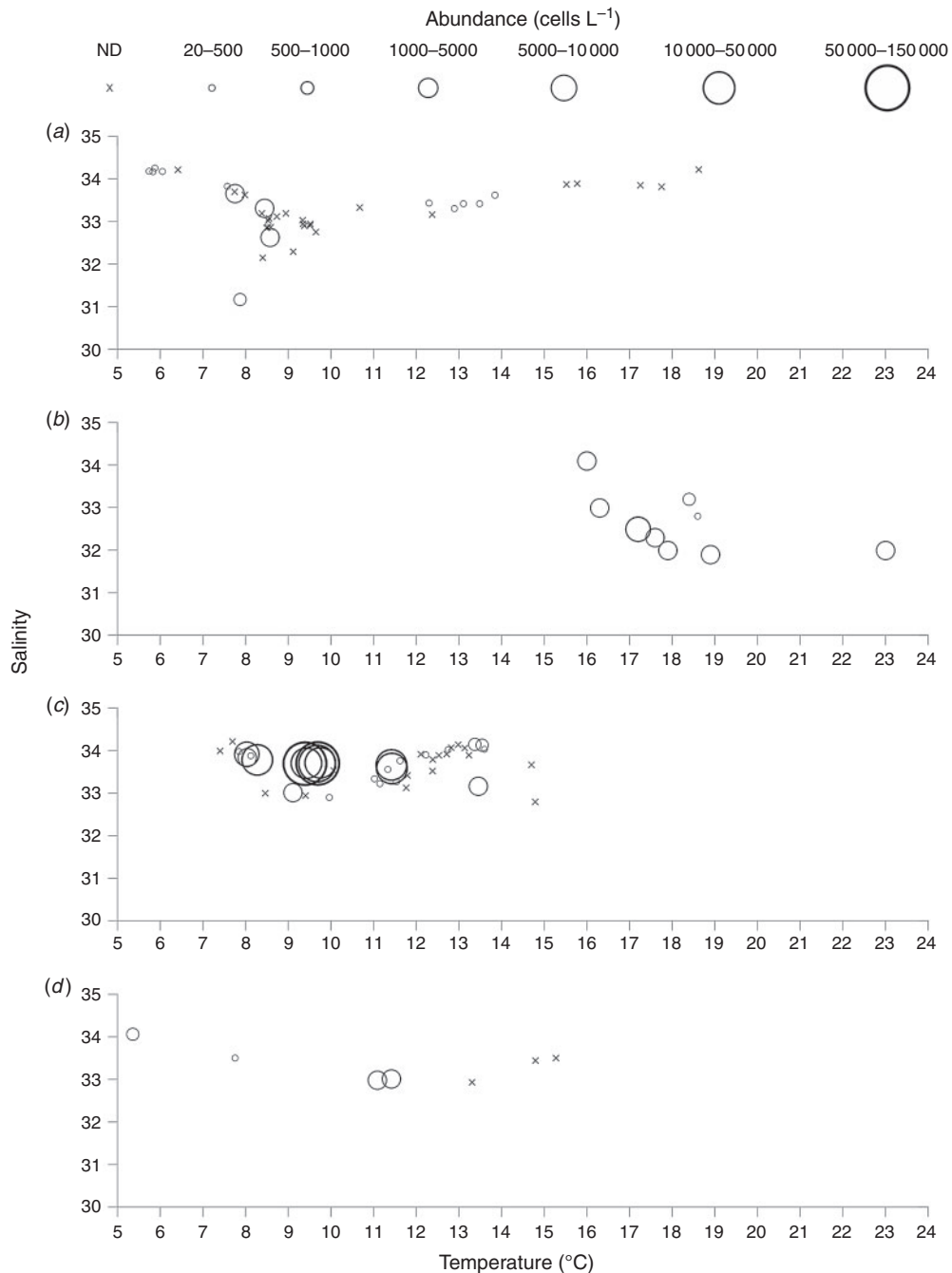
Based on various sampling cruises, Akselman *et al.* (2014) determined that *Azadinium*, albeit at rather low abundances ( $<1 \times 10^3$  cells L<sup>-1</sup>), has a very wide distribution in Argentine waters encompassing northern Argentina and southern Uruguayan shelves, including the mouth of the Río de la Plata (Akselman *et al.* 2014). Similarly, Antacli *et al.* (2018) found unidentified amphidomatacean cells in abundances  $<10^3$  cells L<sup>-1</sup> but widely distributed in the southern Argentine Sea, from ~47 to 53°S. Our results suggest that amphidomatacean species are a regular component of phytoplankton in the area, showing a wide latitudinal distribution from 38 to 55.5°S and covering a wide range of temperatures (5–19°C) and salinities (31–34). Moreover, we confirmed *Azadinium* occurrence in the southern Argentine Sea (~55°S). Maximum *Azadinium*-like cell abundance was found in very diverse plankton assemblages dominated by either diatoms

or dinoflagellates or small phytoflagellates (including haptophytes) and cryptophytes. The highest *Azadinium*-like abundance ( $\sim 1.5 \times 10^5$  cells L<sup>-1</sup>) was recorded in waters with low temperature (9–10°C) and high salinity (33.70; Fig. 7). Similar conditions occurred during the Amphidomataceae blooms from the 1990s (temperatures 7.7–9.2°C and salinity 33.64–33.75; Akselman and Negri 2012), as well as during recent blooms on the external shelf and slope of El Rincón (temperatures 8–10°C; Tillmann *et al.* 2019). By contrast, in Brazilian coastal waters (temperatures 16–22°S), Cavalcante *et al.* (2018) found maximum *Azadinium* spp. (*Az. dexteroporum*, *Az. polongum* and *Az. luciferelloides*) cell densities at temperatures between 21 and 28°C, although these densities (6174 cells L<sup>-1</sup>) were lower than those detected in the present study. Conversely, the occurrence of *Azadinium* in low abundances has been recorded in cold waters ( $<12^\circ\text{C}$ ) from the Northern Hemisphere, in the subpolar area of the Irminger Sea and northern Atlantic Ocean off Iceland (Tillmann *et al.* 2014a), as well as from the Shetland Islands (Tillmann *et al.* 2012b). However, the species found in those studies (*Azadinium trinitatum*, *Az. concinnum*, *Azadinium cuneatum*, *Az. spinosum* and *Az. polongum*) were not detected in the present study. There are limited data on the growth of *Azadinium* based on culture experiments: a North Sea strain of *Az. spinosum* showed a maximum grow rate at 22°C and a considerably lower growth rate at 10°C (Jauffrais *et al.* 2012). An Iceland strain of *Az. polongum* grew well at 10°C, but died rapidly at higher temperatures (Tillmann *et al.* 2012b). Based on these different experimental responses to temperature, it could be expected that *Azadinium* species have distinct species distribution patterns. However, field studies about the spatial and temporal dynamics of *Azadinium* species are still scarce because of their small size and the impossibility of an accurate identification during cell counting under light microscopy. Finally, we cannot rule out that other factors may have a greater effect than temperature on the exceptionally high abundance of Amphidomataceae in the shelf break and El Rincon fronts of the Argentine Sea (e.g. nutrient supply given by the particular oceanographic characteristics of those areas; Acha *et al.* 2004).

Despite the fairly high cell densities of *Azadinium*-like cells reported here, no AZAs were detected in any of the bottle samples. For many samples with a low abundance of *Azadinium*-like cells, this could be due to the chemical detection limit, which, assuming a mean AZA cell quota of 10 fg cell<sup>-1</sup> (Tillmann *et al.* 2014b), would require minimum densities of 2600–44 400 cells L<sup>-1</sup> (depending on detection limits for each expedition) of a toxigenic species to be detected. A lack of AZAs in the high-abundance samples indicates that the dominant species in these populations do not produce any of the AZA congeners analysed. In fact, from 13 species of Amphidomataceae tested so far, AZA production has been proven for only four (Tillmann *et al.* 2009; Krock *et al.* 2012; Rossi *et al.* 2017). However, it is important to consider that *Az. luciferelloides* and *Az. asperum* were described based on Argentinean field samples (Tillmann and Akselman 2016; Tillmann 2018) and thus, because of a lack of cultures, their AZA production potential has not yet been tested.

The toxinological analysis of concentrated plankton net samples obtained during E3 showed the presence of AZA-2 from field plankton samples for the first time in the region.





**Fig. 7.** Abundance of *Azadinium*-like cells at different surface salinities and temperatures for the four expeditions across the Argentine Sea: (a) Expedition 1, (b) Expedition 2, (c) Expedition 3 and (d) Expedition 4.

Despite the presence of a few *Azadinium* cells in these plankton net samples, it is assumed that the large majority of toxin was transferred to predators of *Azadinium* and other protistan grazers. AZA transmission through the food web is known for heterotrophic dinoflagellates and ciliates (Yasumoto 2001; James *et al.* 2003; Krock *et al.* 2008). Moreover, planktonic grazing on *Azadinium* has been suggested as an important loss factor during *Azadinium* blooms (Akselman and Negri 2012).

In the present study, potential AZA vectors, such as aloricate ciliates, tintinnids and heterotrophic dinoflagellates of the genera *Protoperdinium* and *Gyrodinium*, were detected as frequent components of microplankton during E3. Moreover, maximum *Azadinium*-like abundance co-occurred with the maximum density of *G. fusus*, a suspected predator of *Azadinium* sp. (Akselman and Negri 2012), in a sample where AZA-2 was also detected.

Of the 21 AZA analogues analysed in this study, only AZA-2 was detected; the same analogue was exclusively found by Turner and Goya (2015) in mussels collected in Buenos Aires Province (although those authors only analysed AZA analogues 1–3). AZA-2 is known to be produced by *Az. spinosum*, *Az. poporum* and *Am. languida*. Of those, until recently only *Az. poporum*, and here including all Argentinean *Az. poporum* strains (Tillmann *et al.* 2016), was known to produce only AZA-2, whereas strains of *Az. spinosum* were characterised as producing AZA-1 together with AZA-2, and *Am. languida* was characterised as producing AZA-43 together with AZA-2. However, one of many *Az. spinosum* strains recently isolated from the Argentinean shelf was shown to represent the new ribotype B and to produce only AZA-2 (Tillmann *et al.* 2019). Thus, the presence of just AZA-2 in plankton samples (present study) and in Argentinean shellfish (Turner and Goya 2015) perfectly fits the toxin profile of Argentinean *Azadinium* strains available so far, but it is not possible to assign the presence of AZA-2 exclusively to *Az. poporum* or *Az. spinosum*.

## Conclusions

This paper presents the first record of AZA-2 in plankton samples from Argentinean shelf waters, and the first detection of *Az. obesum* in the Southern Hemisphere in addition to four other *Azadinium* species, of which two are potentially toxigenic. Considering that the Atlantic coast of Argentina is a very important region for fisheries and seafood production (Bogazzi *et al.* 2005), the diversity, wide latitudinal distribution and high abundances of *Azadinium*, together with the detection of AZA-2, highlights the potential risk of AZA shellfish contamination episodes. There is a need for more toxinological information and spatiotemporal distribution data, as well as identification of possible source species of AZA in the study area.

## Conflicts of interest

The authors declare that they have no conflicts of interest.

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