

## Influence of re-flooding on phytoplankton assemblages in a temperate wetland following prolonged drought

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### ABSTRACT

Wetlands can experience drying and flooding cycles which influence the dynamics of the phytoplankton assemblages. The aim of our study was to evaluate changes in the phytoplankton structure during a drought/flood period in a warm-temperate wetland. We hypothesized that fluctuations in water level and development of macrophytes favour the development of fast-growing algae with adaptations to low light conditions. We studied algal and cyanobacterial colonization and succession in the nascent planktonic habitat in a wetland in the Southern Hemisphere (Argentina). We assessed changes in phytoplankton biovolume, chlorophyll a concentration (Chl a), richness, diversity, and evenness throughout a drought/flood period. Phytoplankton species were classified into ecological functional groups (FG). Multivariate analysis (RDA) showed that water level, conductivity and percentage macrophyte cover of the site surface (PCSS) explained the variability in the phytoplankton assemblage structure in terms of classes and FG. Particularly, FGs T and L<sub>M</sub> responded to the changes during the drought/flood cycle, probably due to light constraints and stability of the water column induced by the development of emergent and free-floating macrophytes. Our study expands the knowledge of phytoplankton species composition and ecological FG succession under free-floating macrophyte cover in a re-flooding episode. We conclude that water depth and development of macrophytes are the key factors in shaping phytoplankton species structure in a temporary wetland.

Key words: phytoplankton, wetland, shallow lake, drought, flood, biodiversity.

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### INTRODUCTION

There are many types of wetlands worldwide and the heterogeneity of their environments is widely known (Rojo *et al.*, 2012). While sharing common characteristics that define them, these shallow aquatic ecosystems are naturally subjected to water-level fluctuations due to the seasonal variation of precipitation, runoff, and/or evaporation (Naselli-Flores and Barone, 2005). The hydrological variation represents an important factor in shaping wetlands. Climate change models predict increased drought episodes for many regions of the world (Jentsch *et al.*, 2007; IPCC, 2010) as well as an enhanced frequency of extreme hydrometeorological events (Timmermann *et al.*, 1999). Wetlands are amongst the most sensitive ecosystems (Jeppesen *et al.*, 2007; Meerhoff *et al.*, 2007) and may become fragmented during drought periods (Angeler *et al.*, 2010). Recently the number of studies exploring the effects of drought on wetland communities is increasing (Bucak *et al.*, 2012; Granado and Henry, 2014; Rojo *et al.*, 2012; Stević *et al.*, 2013). However, studies are still limited compared to those regarding floods and other disturbances (Lake, 2011).

Algae are key players in the physical, chemical, and biological processes that characterize most freshwater wetlands and their dynamics and composition are influenced by environmental changes triggered by water level fluctuations (Naselli-Flores and Barone, 2005, 2012). In particular, recent studies highlighted that drought events and the associated desiccation are considered as relevant factors controlling the distribution and diversity of algae in freshwater wetlands (Neustupa *et al.*, 2011 and references therein). In this respect, some localities in subtropical wetland systems (Florida, USA) showed different species compositions as a result of differences in the length of the annual dry period; the longer dry period sites are better able to cope with non-periodic events of severe drought (Gottlieb *et al.*, 2005). Goldsborough and Robinson (1996) proposed a conceptual model for wetland algae that includes four quasi-stable states: *dry state*, *open state*, *sheltered state* and *lake state*. The *dry state* is recognized and characterized by the dominance of benthic algae inhabiting the illuminated surface layers of exposed mud flats and submerged sediments. The resting stages of many aquatic organisms persist in the sediments of dry wetlands (Skinner *et al.*,

2001; and references therein), and their abundance and diversity reflects the capacity of an incipient wetland community to respond to re-flooding by producing colonists and resetting the processes of succession (De Stasio, 1990). Then, following flooding, dry wetlands are transformed to a state dependent on the quantity and timing of water input. If the water column is sufficiently shallow so that aquatic macrophytes can develop, the increase in substratum availability may lead to an *open state* wetland in which epiphytes are the predominant algal assemblage. *Sheltered state* is also reached at medium hydrometric levels, but it is characterized by the development of edge-vegetation, which stabilizes the water column, and the dominance of metaphyton. If water input is too rapid for macrophytes to colonize, or if vegetation is excluded by excessively deep water, phytoplankton predominates in a *lake state* wetland (Goldsborough and Robinson, 1996).

Plankton may be especially suitable for studying adaptive processes to environmental change at different spatial and temporal scales, because of the short generation times of many species (Leibold and Norberg, 2004). Because phytoplankton is dispersed easily, air-borne colonization or dispersal through surface connections, and also colonization through propagule banks, could mediate the strength and magnitude of local adaptation (Angeler *et al.*, 2010). It is recognized that the phytoplankton species composition in freshwater ecosystems is influenced by the presence of aquatic macrophytes as they affect the water quality through their effects on nutrient competition, allelopathy, shading and, in the case of floating plants, anoxygenic conditions (Scheffer *et al.*, 2003; Izaguirre *et al.*, 2010; Naselli-Flores and Barone, 2012). Thus, hydrology-driven changes on macrophyte assemblages are expected to further impact on phytoplankton succession.

The phytoplankton community is composed of different functional associations of species that will be favoured or excluded as regards their adaptations to withstand the environmental conditions (Reynolds *et al.*, 2002). In this sense, the functional-group model (Reynolds *et al.*, 2002, and reviewed by Padisák *et al.*, 2009) suggests the selection or exclusion of these adaptive features in the phytoplankton community, which are not specific to a determined phylogenetic group. Thus, the probability of the prevalence of certain functional groups, rather than species dominance, can be better narrowed down for a given set of environmental conditions. The functional group approach helps to understand why differing phytoplankton species are selected over others in an assembling community, as well as why its composition varies at both a temporal and spatial scale (Reynolds *et al.*, 2002; Naselli-Flores and Barone, 2012; Stević *et al.*, 2013).

In this work we aim to analyze the changes in phytoplankton structure and dynamics in a warm-temperate wetland starting from a *dry state* during a complete annual

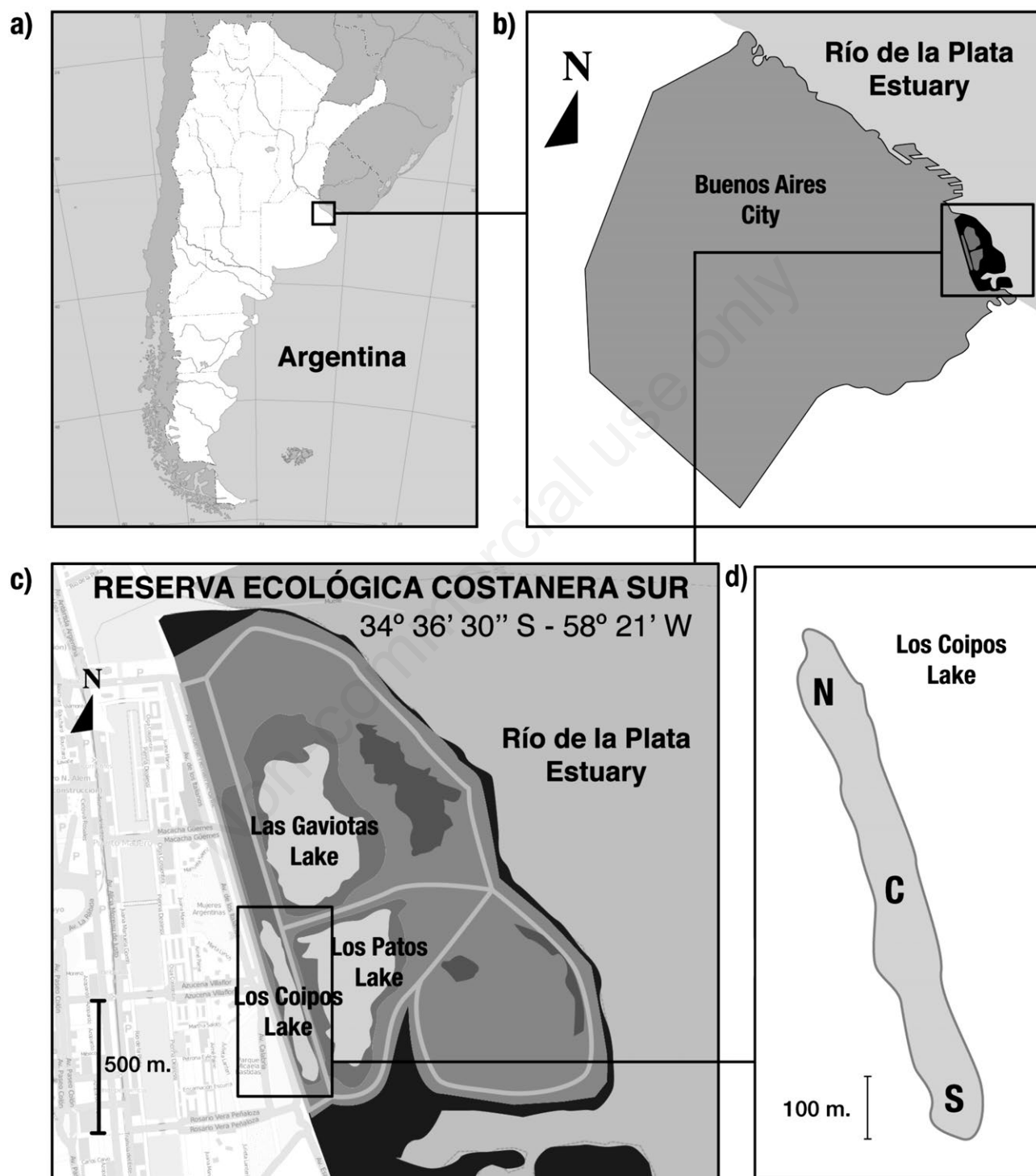
drought/flood cycle. We explore the response of the phytoplankton species composition and functional-group assemblage to the re-flooding of the basin. We hypothesize that rapid changes in water level and macrophyte cover (emergent and/or free floating) affect the phytoplankton composition and favour functional groups represented by small fast-growing algae with adaptations to low light conditions. As well, the persistence of a free-floating macrophyte cover favours algae with adaptations to withstand the physical stability of the water column imposed by the presence of macrophytes, and the low light conditions encountered beneath the coverage.

## METHODS

### Study site

The *Reserva Ecológica Costanera Sur* (RECS hereafter) (34° 36'S - 58° 21'W) is located on the east of Buenos Aires city (Argentina) (Fig. 1). This area was created in 1978, by reclaiming land from the river; by ground-filling with sediments from the Rio de la Plata. Three shallow water bodies were created within the reserve, delimited by constructed levees without river connection. These management efforts continued until 1984 when it was declared a natural park and ecological reserve in 1986. Despite its anthropogenic origin, RECS can be considered a wetland following the criteria of the Wetland Convention (Davis *et al.*, 1996) and since 2005, the area was included in the list of *Wetlands of International Importance* of the International Convention on Wetlands. This wetland is in the temperate region of the Southern Hemisphere, with mean annual precipitation of 800-1000 mm (Servicio Meteorológico Nacional, 2012). In recent years, the area has been influenced by increased frequency of extreme hydrometeorological events such as El Niño Southern Oscillation (ENSO), measured by the Southern Oscillation Index (SOI). When the SOI is strongly negative (El Niño) intense rains and mayor flooding may occur in this area and, contrarily, when SOI is positive (La Niña) severe droughts may occur.

In this work, we studied one of the shallow lakes of RECS, Los Coipos Lake. Since it was built until 2008, the water level of this shallow lake has fluctuated seasonally, generally increasing in autumn (maximum depth: 1.5 m) and decreasing in summer (Zapata, *personal communication*). The summer season is critical due to the high temperatures that increase evaporation and that can result in drying out of the lake. The hydrological behavior of recent years characterizes Los Coipos shallow lake as a temporary water body that alternates between drought and flooded states. This shallow lake was completely dry in summer 2008, its watershed was colonized by grassland until autumn 2009 when it was subjected to re-flooding. During 2007-2008 the region was under a moderate effect



**Fig. 1.** a) Republic of Argentina map. b) Buenos Aires city map. c) Reserva Ecológica Costanera Sur (RECS) scheme, with Los Coipos Lake on the box. Modified from [http://www.buenosaires.gov.ar/areas/med\\_ambiente/reserva/mapa.php](http://www.buenosaires.gov.ar/areas/med_ambiente/reserva/mapa.php). d) Los Coipos Lake indicating the sampling sites: N, site North; C, site Central; S, site South.

of positive SOI (La Niña) and in April 2009 a strong negative SOI (El Niño) effect started. The study was carried out throughout this *El Niño* period from August 2009, when the basin of the shallow lake started to be refilled, until August 2010 when it ended-up completely covered by a dense macrophyte cover.

### Sampling methodology

To test our hypotheses typical limnological variables (both biotic and abiotic) were measured during a drought/flood period (August 2009-August 2010) starting from a *dry state* (Goldsborough and Robinson, 1996). Water samples for physical, chemical and biological analyses were collected from Los Coipos Lake at three sites: South (S), Central (C) and North (N), coinciding with previously studied sites (Fazio and O'Farrell, 2005). Each of the three sites was sampled on 14 occasions, resulting in a total of 42 samples. Initially, three isolated ponds located within the watershed of the dried shallow lake were sampled fortnightly (*initial period* hereafter: 19-Aug-09, 2-Sep-09, 15-Sep-09); and then, when the three ponds connected together into a single and continuous water body, approximately at month intervals (*re-flooding period* hereafter: 28-Sep-09, 19-Oct-09, 9-Nov-09, 30-Nov-09, 21-Dec-09, 25-Jan-10, 8-Mar-10, 5-Apr-10, 3-May-10, 29-Jun-10, 19-Aug-10).

At all sampling dates we recorded the vegetation type (if present) in each of the three sampling sites. We also estimated the percentage cover of the site surface (PCSS) by emergent and free floating macrophytes using four categories: no vegetation, up to 25%, 26- 75%, and 76-100%. To study the evolution of the shallow lake during the drought/flood interval 2009-2010, Landsat 5TM images (Path 225, Row 84) that were available for the period were used. We obtained four images characteristic of four well defined scenarios: February 4, 2009 (initial *dry state*) which persisted until the re-flooding on August 2010; October 2, 2009 (incipient formation of a continuous water body); January 22, 2010 (sharp water level decrease); September 19, 2010 (shallow lake with complete surface coverage by emergent and floating macrophytes). A sub-area covering the study site was extracted and geometric correction was performed using well distributed Ground Control Points, with a positional accuracy of  $\pm 1$  pixels. A radiometric correction was conducted with the Rayleigh dispersion model (Stumpf, 1992). The total lake area was determined as the largest area found during high waters. In order to estimate the percent cover of macrophytes in the lake an unsupervised classification was performed based on ISODATA clustering with a variability threshold of 98% in each image. We used the signature editor in ERDAS Imagine to extract spectral data from the image for the classes (vegetation and water). The classification of the satellite data and the analysis of spectral signatures together with the *in situ*

observations allowed us to characterize the scenarios at the whole lake level. Image processing was conducted using ERDAS Imagine 9.1 software.

### Physical and chemical analyses

Dissolved oxygen (DO) concentration, water temperature, pH, and conductivity were measured *in situ* with Hanna HI8314 and Hanna HI8033 portable electronic sensors (Hanna Instruments, Smithfield, RI, USA) and the transparency was estimated using a Secchi disk. The water depth of each site was measured with a meter stick. Water samples were collected for the determination of the main dissolved nutrients for algal and cyanobacterial growth; soluble reactive phosphorus (SRP), ammonium (N-NH<sub>4</sub>) and nitrates+nitrites (N-NO<sub>3</sub>+N-NO<sub>2</sub>). The concentrations of these nutrients were determined using a HACH® DR/2010 (HACH® Company, USA) spectrophotometer and the corresponding kits of HACH® reagents (detection limits: 0.001-2.500 mg L<sup>-1</sup> for SRP; 0.001-2.500 mg L<sup>-1</sup> for N-NH<sub>4</sub>; 0.01-0.50 mg L<sup>-1</sup> (low range) and 0.3-30.0 mg L<sup>-1</sup> (high range) for N-NO<sub>3</sub>+N-NO<sub>2</sub>). Samples for nutrient analysis were filtered through Whatman® GF/F filters. SRP concentration was estimated by the ascorbic acid method, the concentration of N-NH<sub>4</sub> was determined by the indophenol-blue method, and that of N-NO<sub>3</sub>+N-NO<sub>2</sub> was estimated by the cadmium reduction method (APHA, 2005). Dissolved inorganic nitrogen (DIN) represents the sum of N-NH<sub>4</sub>+N-NO<sub>3</sub>+N-NO<sub>2</sub>. In addition, water samples were filtered with Whatman® GF/C in the laboratory in order to determine the concentration of suspended solids (SS) by weighing the dry residue at 103° C (APHA, 2005).

### Biological analyses

Samples for the estimation of phytoplankton chlorophyll *a* concentration (Chl *a*), and for the qualitative and quantitative analyses of phytoplankton were obtained in PVC flasks. Chl *a* was estimated spectrophotometrically using hot ethanol (60-70°C) extraction (Lorenzen, 1967; Nusch, 1980). Qualitative cyanobacterial and algal samples were obtained with a 15 µm pore net, a sub-sample was fixed in 4% formalin and a second sub-sample was used for *in vivo* taxonomical identification using a light microscope (Zeiss®) at X1000 magnification and up-to date phycological references such as Komárek and Anagnostidis (1999), Komárek and Anagnostidis (2005) for cyanobacteria; Hüber-Pestalozzi (1941) and Starmach (1985) for Chrysophyceae; Javornicky (2003) for Cryptophyceae; Krammer and Lange-Bertalot (1986, 1988, 1991a,b), Lange-Bertalot (2001), Metzeltin and García-Rodríguez (2003) and Zalocar de Domitrovic and Maidana (1997) for Bacillariophyceae; Komárek and Fott (1983) for Chlorococcales; Prescott *et al.* (1975, 1977, 1981) for Zygnematophyceae; Ettl (1978) for Tribon-

phyceae; Tell and Conforti (1986) for Euglenophyceae.

Quantitative ( $>2 \mu\text{m}$ ) phytoplankton samples were fixed with 1% acidified Lugol's iodine, and those for the quantification of picophytoplankton ( $0.2\text{-}2 \mu\text{m}$ ) were preserved with 2% ice-cold glutaraldehyde. In order to estimate the abundance of each of the phytoplankton size-fractions, two sub-samples were counted and averaged for each sampling date and site. The counts of the  $>2 \mu\text{m}$  phytoplankton fraction were carried out to species level whenever possible, using an inverted microscope (Zeiss®) following the Utermöhl (1958) technique at X400 magnification on chambers of 5 or 10 cc after at least 24 h sedimentation. Counting errors were estimated according to Venrick (1978), accepting a maximum of 20% for the most abundant species. Algae were classified in the differing classes and according to the functional groups (FG) proposed by Reynolds *et al.* (2002) and reviewed by Padisák *et al.* (2009). Biovolumes of all algal species were calculated following Hillebrand *et al.* (1999) and Sun and Liu (2003). Then, considering the abundances and the calculated biovolume for each species, we estimated the total biovolumes of each of the algal classes and cyanobacteria, as well as those of each of the FG.

Picophytoplankton fraction was counted from the fluorescence given off by photosynthetic pigments following Kemp *et al.* (1999); 2 mL sub-samples were filtered through  $0.2 \mu\text{m}$  pore sized black polycarbonate filters. Each filter was examined for pigment autofluorescence with a Zeiss® Axioplan microscope equipped with an HBO 50W lamp, and a filter set for blue light excitation (BP 450-490 nm, FT 510 nm, LP 520 nm) and green light excitation (BP 546 nm, FT 580 nm, LP 590 nm) were used in order to identify picoeukaryotic algae and picocyanobacteria, respectively. A minimum of 400 cells were counted (corresponding to at least 20 fields of view) at X1000 magnification, accepting a maximum of 15% error.

### Data analyses

We estimated different attributes of phytoplankton assemblage: species richness, diversity (Shannon-Wiener) (Magurran, 1988) and evenness (Pielou, 1966) for each sample site and date (a total of 42 samples) in terms of biovolume. Total species richness of Los Coipos Lake was calculated as total number of species encountered in water column, considering both the qualitative and quantitative study of  $>2 \mu\text{m}$  phytoplankton, throughout the study period. Average species richness, diversity and evenness were calculated as the mean value of the three sites (S, C and N) per sampling date for each variable.

We performed descriptive statistical analysis of the physical, chemical and biological variables (SPSS Statistics 17.0 – StatSoft, USA). Non-parametric correlations were performed with Spearman's Rho coefficient among the abiotic variables, abiotic parameters vs algal classes, abiotic

variables vs biovolumes of algal classes or biovolumes of FG. We calculated Stander's similarity index SIMI (Elber and Schanz, 1989), in order to compare, in pairs, phytoplankton community structure of different sites in temporal succession using abundance values. This index ranges between 0-1, with 1 representing the highest similarity between a pair of communities. Direct ordination analyses were used to assess for significant relationships between environmental variables and phytoplankton species biovolume, including species that represented more than 5% of the total  $>2 \mu\text{m}$  phytoplankton biovolume. Previously, we performed a Detrended Correspondence Analyses (DCA) and, as data showed a linear response, we applied a Redundancy Analysis (RDA) in order to select the minimum number of variables that explained the highest variability in phytoplankton composition. The biological data were log-transformed. All environmental parameters that were not highly correlated ( $r > 0.90$ ) were included in the analysis: water depth, conductivity, pH, DIN, SRP, PCSS, presence/absence of emergent macrophytes (EM), and presence/absence of free-floating macrophytes (FFM). PCSS was considered as a categorical variable: 0, no vegetation; 1, up to 25%; 2, 26-75%; 3, 76-100%. Significance of ordination axes was assessed by Monte Carlo permutation test (499 permutations). Relative contribution of each of environmental variables to the ordination axes was evaluated by intraset correlations. Multivariate analyses were performed using CANOCO program for Windows 4.5 (Canonical Community Ordination) (ter Braak and Smilauer, 2002).

### RESULTS

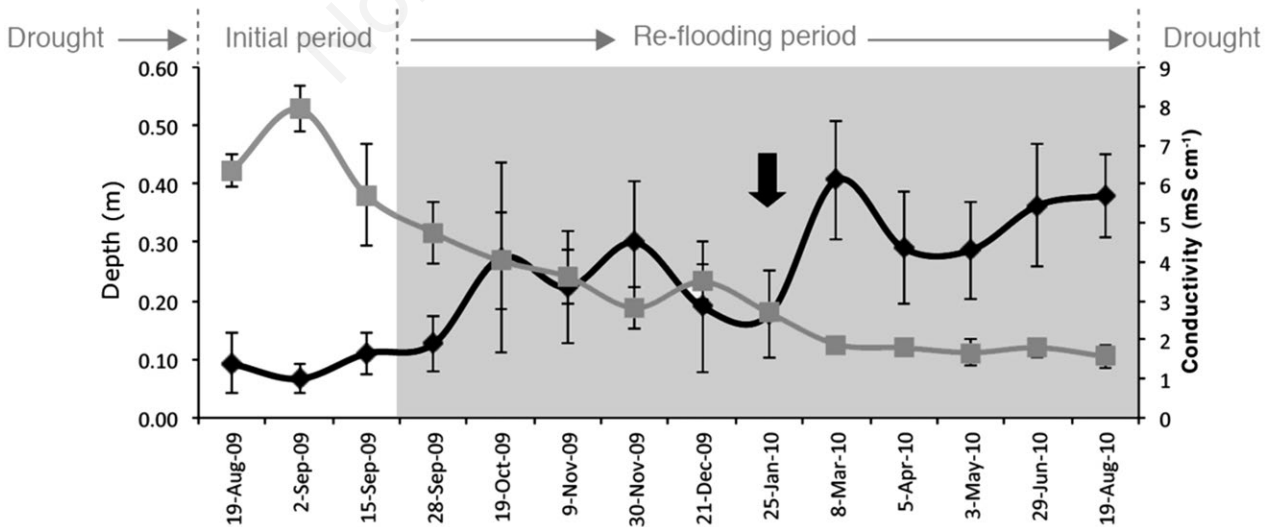
Los Coipos Lake was characterized by wide water-level fluctuations from its initial *dry state* (December 2008-August 2009) and as it was subjected to re-flooding during the drought/flood period 2009-2010. In February 4 2009, the satellite image (Fig. 2a) and the spectral signatures of classes derived from the classification technique showed that the entire lake area was covered by vegetation (reflectance peaks at green and near infrared bands), mostly comprised by grassland that colonized when the lake had no surface water. This image coincides with the dry period that occurred before our study. With the onset of re-flooding in August 2009, the lake gradually changed: on October 2, 2009 (Fig. 2b), 55% of the lake area, comprising mainly the central region, was represented by open-water (low reflectance in visible region of the spectra and no reflectance in near and far infrared regions). Interestingly, within this area, aquatic vegetation cover (*Typha latifolia*) tended to increase towards the littoral zone. Moreover, the edge of the lake was covered by dry vegetation, as depicted by the higher reflectance in the far infrared region of the spectral signature. This dry vegetation represents the remnant grassland, which de-

cayed with the increase in water level and was gradually replaced by aquatic plants. In January 22, 2010 (Fig. 2c), the central open-water region comprised only 18% of lake area, as the littoral aquatic vegetated area increased to 63%, including *Azolla filiculoides* and *Pistia striatotes*. In early summer (December 2009-January 2010) the lake decreased in its water depth (Fig. 3) due to high water

temperatures and low precipitation measured in that summer period. During this period, the increased concentration of Chl *a* in the water column was evidenced by lower reflectance in the blue region and a small peak in the near infrared region of the spectra. Both northern and eastern edges were still covered by dry grassland vegetation. Regarding the final state, in September 19, 2010 (Fig. 2d)



**Fig. 2.** Thematic maps representing the surface coverage of the different landscape elements from Los Coipos Lake obtained from Landsat 5TM images available for the study period. N, site North; C, site Central; S, site South. a) Feb 2009. b) Oct 2009. c) Jan 2010. d) Sep 2010.



**Fig. 3.** Average temporal fluctuations of water depth (black) and conductivity (grey) in Los Coipos Lake during the study period (August 2009-August 2010). The shaded background indicates the formation of a continuous shallow lake. The bars indicate  $\pm 1$  standard deviation.

no open-water area was detected for the lake surface; spectral signatures denote that 100% of the area was covered by aquatic vegetation. Particularly, the central zone was covered by free-floating plants mainly *Azolla filiculoides* (greater reflectance peaks in red and near infrared regions), whereas littoral areas were covered by both free-floating and emergent macrophytes (*Azolla filiculoides*, *Pistia striatotes* and *Typha latifolia*).

### Physical and chemical properties

Ours results showed marked differences in physical and chemical characteristics of the three studied sites in Los Coipos Lake throughout an annual drought/flood period (Tab. 1). Initially, the three sites consisted of isolated pools, with depths not exceeding 0.15 m (Fig. 3) until early spring (four weeks later) when the three pools connected together into a single shallow water body. Throughout the study, water depth was similar at all three sites. Conductivity was high (8.56 mS cm<sup>-1</sup>) at the beginning of the study (August-September 2009) and decreased with increasing water depth (Fig. 3). Average pH values (Tab. 1) were minimum in winter (June 2010) and maximum in the summer (January 2010). A positive correlation between this variable and temperature (n=42, r=0.54, P<0.0001) was encountered. DO concentrations (Tab. 1) showed large fluctuations in temporal scale, which did not respond to a seasonal pattern. Minimum values were recorded when the shallow lake surface was covered by free-floating macrophytes (April-August 2010). We found a positive correlation between this variable and pH (n=42; r=0.52; P<0.0001). The SS concentrations (Tab. 1) were variable throughout the study; the highest levels were found when the water depth was low. In the absence of free-floating macrophytes, Secchi disk depth reached the bottom of the lake.

The concentration of dissolved nutrients (SRP, N-NO<sub>3</sub>+N-NO<sub>2</sub> and N-NH<sub>4</sub>) showed differences among

the sites during the study period, presenting a relatively wide range of values (Tab. 1). The highest concentrations of N-NH<sub>4</sub> values were found when the lake surface was totally covered with floating macrophytes. DIN ranged from 0.02 to 1.30 mg L<sup>-1</sup>, and the lowest values were found at the beginning and at the end of the study period. The proportions of N-NH<sub>4</sub> / N-NO<sub>3</sub>+N-NO<sub>2</sub> were variable and at the end of the study the ratio was highest.

### Algal assemblage attributes

A high positive correlation between abundance and biovolume (n=42, r=0.87, P<0.0001) was encountered for all algal classes, except Chlorophyceae (n=42, r=0.56, P<0.0001) and cyanobacteria (n=42, r=0.74, P<0.0001). Fluctuations were detected in picophytoplankton, >2 μm phytoplankton species diversity, species richness and evenness of phytoplankton assemblage of Los Coipos Lake throughout the drought/flood period 2009-2010 (Tab. 2). Phytoplankton Chl *a* concentration and >2 μm phytoplankton showed marked variations (Fig. 4). The highest values of Chl *a* did not always coincide with the >2 μm phytoplankton biovolume. The picophytoplankton composition consisted of phycocyanin-rich picocyanobacteria (0.97-5.3 mm<sup>3</sup> L<sup>-1</sup>) that dominated the picoeukaryotic algae (<1.2 mm<sup>3</sup> L<sup>-1</sup>) in all samples. Within the larger phytoplankton >2 μm fraction, we recorded a total of 191 algal species and 44 cyanobacteria when adding up all the taxa encountered in the qualitative and the quantitative analyses. The floristic list of our study shows 235 species (Supplementary Tab. 1). Of those, 140 species were present in the quantitative samples belonging to eight algal classes and cyanobacteria. Algal species richness was similar in the three sites and the lowest values were recorded at the beginning and the end of this study (Tab. 2). Minimum evenness was found at the beginning of the study (0.14 on site C in August 2009), and the maximum was recorded with the formation of the lake (0.82 on Novem-

**Tab. 1.** Descriptive statistics of the physical and chemical parameters measured in Los Coipos Lake (annual averages) during the study period (August 2009-August 2010). Mean and range (min-max).

Variable	N	Site S	Site C	Site N
Depth (m)	14	0.18 (0.07-0.32)	0.33 (0.09-0.52)	0.20 (0.04-0.38)
Conductivity (mS cm <sup>-1</sup> )	14	3.73 (1.30-7.40)	3.79 (1.52-7.79)	3.21 (1.43-8.56)
Temperature (°C)	14	18.40 (10.40-28.00)	17.84 (8.70-27.00)	16.72 (7.20-27.50)
pH	14	7.52 (6.03-9.57)	7.89 (5.23-10.45)	7.90 (4.92-9.83)
DO (ppm)	14	7.80 (1.64-15.51)	7.20 (1.35-12.99)	6.68 (0.22-11.71)
SS (mg L <sup>-1</sup> )	14	26.38 (8.50-49)	14.24 (4.00-29.00)	53.48 (6.00- 62.80)
SRP (mg L <sup>-1</sup> )	14	0.09 (0.02-0.18)	0.07 (0.03-0.11)	0.12 (0.03-0.24)
N-NH <sub>4</sub> (mg L <sup>-1</sup> )		0.04 (0-0.10) (N=14)	0.03 (nd-0.09) (N=13)	0.07 (0-0.21) (N=14)
N-NO <sub>3</sub> +N-NO <sub>2</sub> (mg L <sup>-1</sup> )		0.49 (0.01-1.30) (N=14)	0.51 (nd-0.80) (N=13)	0.34 (nd-0.70) (N=11)
DIN (mg L <sup>-1</sup> )	14	0.56 (0.20-1.30)	0.50 (0.03-0.83)	0.41 (0.02-0.91)

N, number of samples; Site S, site South; Site C, site Central; Site N, site North; DO, dissolved oxygen concentration; SS, suspended solids; SRP, soluble reactive phosphorus; NH<sub>4</sub>, ammonium; NO<sub>3</sub>+NO<sub>2</sub>, nitrates+nitrites; DIN, dissolved inorganic nitrogen; nd, not detectable.

ber 2009) and with the development of a profuse free-floating macrophyte cover (0.80 on June 2010). Diversity (Shannon-Wiener) values ranged between 0.34 and 2.63. The characteristic habitat of the cyanobacteria and algae encountered in the water column fluctuated according to water depth of the lake (Fig. 5 a,b,c). We observed a relative high algal biovolume, mainly represented by benthic species, during the *initial period* when water depth was low and the lake comprised isolated ponds. The relative importance of benthic algae decreased with increasing depths. In the beginning of the *re-flooding period* (late September-December 2009) we observed an enhanced development of epiphytic algae and cyanobacteria. The relative contribution of different algal classes and cyanobacteria to total >2 µm phytoplankton biovolume also fluctuated throughout the studied period (Fig. 5 d,e,f). Initially, the three sites differed in their algal class and cyanobacterial composition being dominated by Euglenophyceae at site S (negative correlation with conductivity:  $n=42$ ,  $r=-0.46$ ,  $P<0.002$ ) (Fig. 5d), Dinophyceae at site C (Fig. 5e) Bacillariophyceae and Chlorophyceae at site N (Fig. 5f). In all three sites, Bacillariophyceae increased their relative abundance until the formation of the continuous shallow lake and then; in the *re-flooding period*, their relative contribution was variable, though a decreasing pattern was observed as the lake was covered with macrophytes (negative correlation with EM emergent macrophytes:  $n=42$ ,  $r=-0.47$ ,  $P<0.002$ ). Chlorophyceae and cyanobacteria were present throughout the study, with variable contributions. A positive correlation between Cyanobacteria and SRP was encountered ( $n=42$ ,  $r=0.44$ ,  $P<0.004$ ). At the end of the study, Euglenophyceae and

Cryptophyceae were co-dominant at site S (Fig. 5d) Chlorophyceae and Cryptophyceae were also co-dominant at site C (Fig. 5e), while Cyanobacteria were dominant at site N (Fig. 5f). Furthermore, Euglenophyceae and Cryptophyceae showed a positive correlation with free-floating macrophytes cover FFM ( $n=42$ ,  $r=0.50$ ,  $P<0.001$  and  $n=42$ ,  $r=0.55$ ,  $P<0.0001$ , respectively).

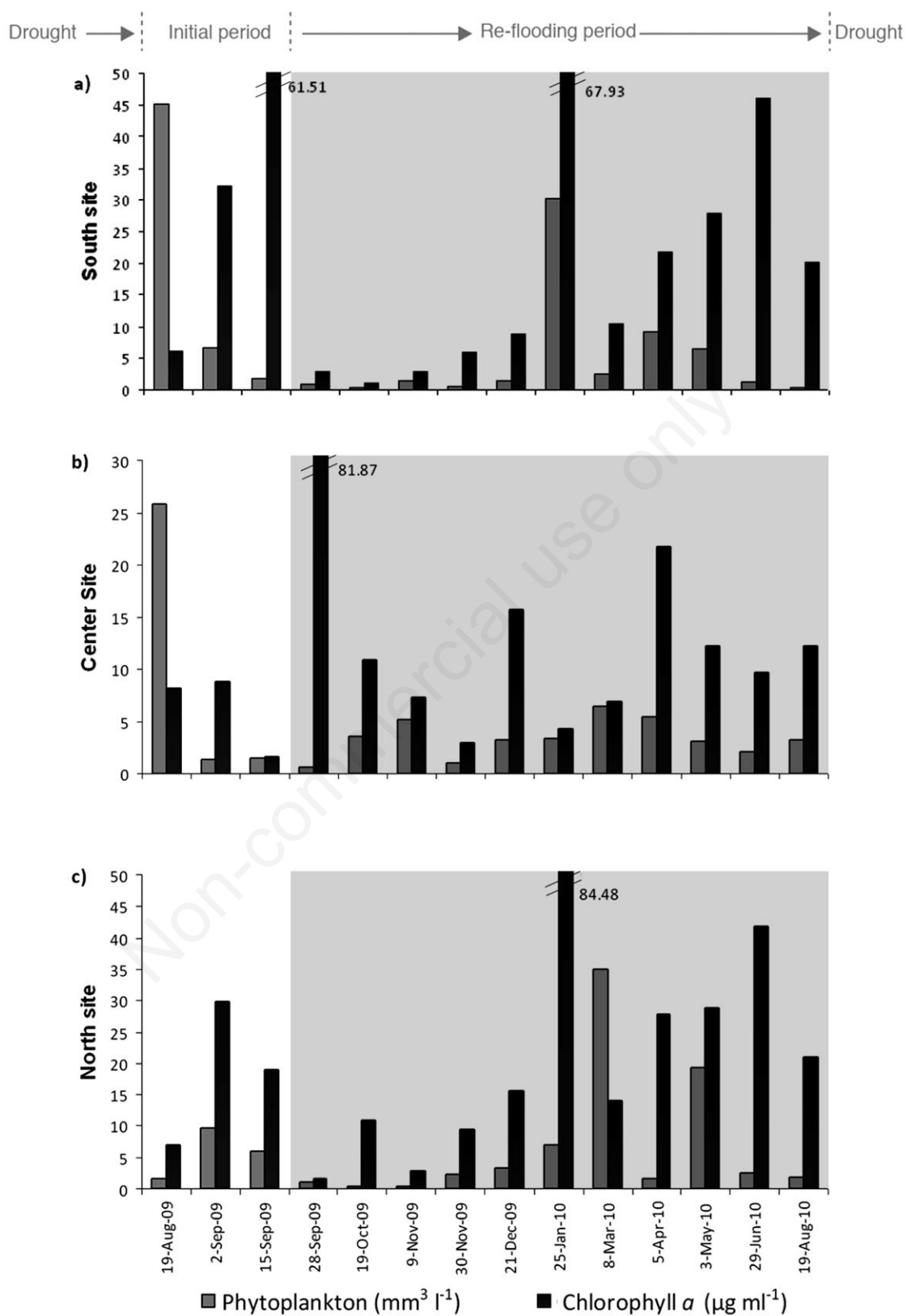
As regards functional group classification, 12 of the 39 FG proposed were encountered in a proportion that exceeded 2% of the total biovolume, with variable dominance throughout the drought/flood cycle (Tab. 3). Initial proportions of FGs of the three sites were different and dominated by W1 (*Euglena caudata*) and Y (*Cryptomonas cf. obovata*); L<sub>O</sub> (*Peridinium* sp.); C and X1 (*Stephanocyclus meneghiniana* and *Chlamydomonas* spp.) for sites S, C and N respectively. In the *re-flooding period*, when the three pools were connected in a continuous water body, we observed a different contribution of FGs to the phytoplankton community. Assemblages mainly represented were W2, Y, D, T, W1, M, L<sub>M</sub> and C. Towards the end of the study period when the whole shallow lake was covered by a dense free-floating plant cover (mainly *Azolla filiculoides*), sites S and C were similar in terms of the proportion of co-dominants W1 and Y, whereas at site N, L<sub>M</sub> was clearly dominant. Significant correlation coefficients were found between D, W1, W2 and Y vs FFM ( $n=42$ ,  $r=0.51$ ,  $P<0.001$ ;  $n=42$ ,  $r=0.43$ ,  $P<0.005$ ;  $n=42$ ,  $r=0.49$ ,  $P<0.001$  and  $n=42$ ,  $r=0.55$ ,  $P<0.0001$  respectively); whereas W1 showed a negative correlation with conductivity ( $n=42$ ,  $r=-0.40$ ,  $P<0.008$ ), W2 with pH ( $n=42$ ,  $r=-0.42$ ,  $P<0.06$ ) and T<sub>B</sub> with water depth ( $n=42$ ,  $r=-0.4$ ,  $P<0.009$ ). Analysis of the similarity of the community structure by means of the SIMI index, revealed

**Tab. 2.** Ranges of biovolume for different phytoplankton attributes recorded in Los Coipos Lake, throughout the study period (August 2009-August 2010). Range (min-max).

Variable	N	Site S	Site C	Site N
Chl a (µg L <sup>-1</sup> )	14	(1.08-67.93)	(1.63-81.87)	(0-84.48) (N=13)
Picophytoplankton (mm <sup>3</sup> L <sup>-1</sup> )	14	(1.48-4.33)	(1.17-4.38)	(1.96-6.55)
Picocyanobacteria (mm <sup>3</sup> L <sup>-1</sup> )	14	(1.25-3.20)	(0.97-4.01)	(1.75-5.33)
Picoeukaryotic algae (mm <sup>3</sup> L <sup>-1</sup> )	14	(0.21-1.13)	(0.14-0.92)	(0.19-1.22)
Phytoplankton >2 µm (mm <sup>3</sup> L <sup>-1</sup> )	14	(0.41-45.20)	(0.57-25.79)	(0.45-35.07)
Cyanobacteria (mm <sup>3</sup> L <sup>-1</sup> )	14	(0-0.63)	(0-0.60)	(0.02-21.59)
Bacillariophyceae (mm <sup>3</sup> L <sup>-1</sup> )	14	(0.08-7.61)	(0.01-1.93)	(0.05-8.97)
Chlorophyceae (mm <sup>3</sup> L <sup>-1</sup> )	14	(0-1.18)	(0.03-5.03)	(0.04-3.85)
Chrysophyceae (mm <sup>3</sup> L <sup>-1</sup> )	14	(0-3.67)	(0-0.15)	(0-0.16)
Cryptophyceae (mm <sup>3</sup> L <sup>-1</sup> )	14	(0-11.38)	(0-2.59)	(0-0.26)
Dinophyceae (mm <sup>3</sup> L <sup>-1</sup> )	14	(0-0.34)	(0-24.02)	Not found
Euglenophyceae (mm <sup>3</sup> L <sup>-1</sup> )	14	(0-30.80)	(0-2.12)	(0-3.38)
Tribophyceae (mm <sup>3</sup> L <sup>-1</sup> )	14	(0-0.29)	Not found	Not found
Zygnematophyceae (mm <sup>3</sup> L <sup>-1</sup> )	14	(0-0.03)	(0-1.76)	(0-0.17)
Species richness	14	(13-42)	(15-42)	(13-45)
Diversity (Shannon-Wiener)	14	(0.85-2.63)	(0.34-2.33)	(0.58-2.22)
Evenness	14	(0.33-0.82)	(0.14-0.71)	(0.17-0.75)

N, number of samples; Site S, site South; Site C, site Central; Site N, site North; Chl a, chlorophyll a.



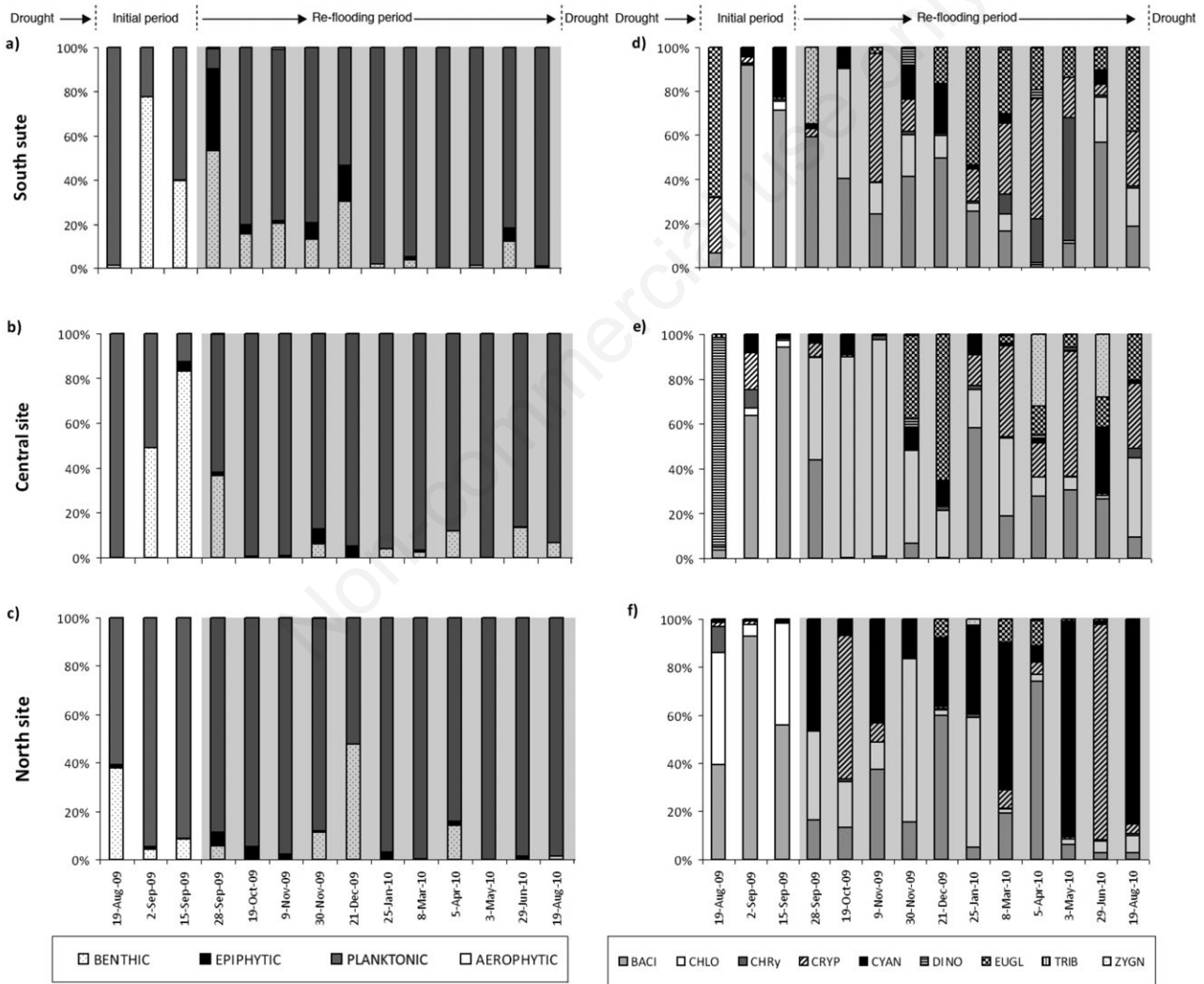


**Fig. 4.** Phytoplankton  $>2 \mu\text{m}$  biovolume (gray bars) and phytoplanktonic Chl *a* concentration (black bars) in Los Coipos Lake during the study period (August 2009-August 2010). a) Site South. b) Site Central. c) Site North. The shaded background indicates the formation of a continuous shallow lake.

that, whereas the three sites showed strong differences initially (site N vs site S: 0.33; site N vs site C: 0.09, and site C vs site S: 0.04), at the end of the study their community structure was notably more similar (site N vs site S: 0.63; site N vs site C: 0.79 and site C vs site S: 0.84).

The eigenvalues of RDA axis 1 (0.14) and axis 2 (0.07) accounted for 21.3% of the cumulative variance in the biovolume of 37 species. The species-environment correlations of RDA axes 1 and 2 were high (0.93 and 0.84, respectively), and the first two axes accounted for 60.8% of the variance in the species-environment relationships. The factor that most affected the algal assem-

blage was the presence of free-floating macrophytes (intra-set correlation coefficient: -0.82) (Fig. 6). The second set of important factors were conductivity, presence of emergent macrophytes, PCSS, and water depth (intra-set correlation coefficients: 0.80, -0.70, -0.68 and -0.67, respectively). Samples belonging to the *initial period* (1, 2, 3) and 4 S, clearly aggregated together in the lower-right side of the triplot characterized by high conductivity values. Most of the samples of the early *re-flooding period* (4 C, 4 N, 5, 6, 7, 8) aggregated in the upper-right side, whereas those of the late *re-flooding period* with further colonization by emergent and free-floating macrophytes



**Fig. 5.** a, b, c) Proportions of the habitats of cyanobacteria and algal classes to >2 μm phytoplankton. d, e, f) Relative contribution of cyanobacteria and algal classes to >2 μm phytoplankton biovolume in Los Coipos Lake water column during the study period (August 2009-August 2010). a, d) Site South. b, e) Site Central. c, f) Site North. The shaded background indicates the formation of a continuous water body. BACI, Bacillariophyceae; CHLO, Chlorophyceae; CHRy, Chrysophyceae; CRYP, Cryptophyceae; CYAN, Cyanobacteria; DINO, Dinophyceae; EUGL, Euglenophyceae; TRIB, Tribophyceae; ZYGN, Zygnematophyceae.

(9, 10, 11, 12) aggregated at the left side of the figure. The final period samples also aggregated together (13, 14) at the upper-left side. During that period, there was a complete macrophyte coverage of the shallow lake. It is worth mentioning that the sample 13 corresponding to the site C (13 C) did not have complete macrophyte coverage of its surface and is discrete from the rest of the samples of the final period.

## DISCUSSION

The phytoplankton species structure and dynamics, as well as the dominant ecological functional groups, of a warm-temperate wetland subjected to drought/flood periods, responded to monthly water level shifts and to macrophyte cover fluctuations during an annual cycle. Starting from a *dry state*, the re-flooding and the gradual colonization by free-floating macrophytes (mainly *Azolla filiculoides*) of Los Coipos Lake, affected the water conditions and, as hypothesized, determined the phytoplankton assemblage in this wetland. The initial water depth registered during the studied drought/flood period was markedly lower than the one registered in previous studies (Fazio and O'Farrell, 2005). This fact is associated with the reduced precipitation in preceding years, resulting from the moderate and strong events of La Niña, which contributed to the drought of the region. May 2009 marked a new wet period that provoked the beginning of the re-flooding of the lake three months later (August 2009) when we initiated the study. In just 4 weeks the basin of the lake was re-filled. From June 2010 onwards, La Niña dominated again yielding to the gradual desiccation of this ecosystem. The effects of the shift between El

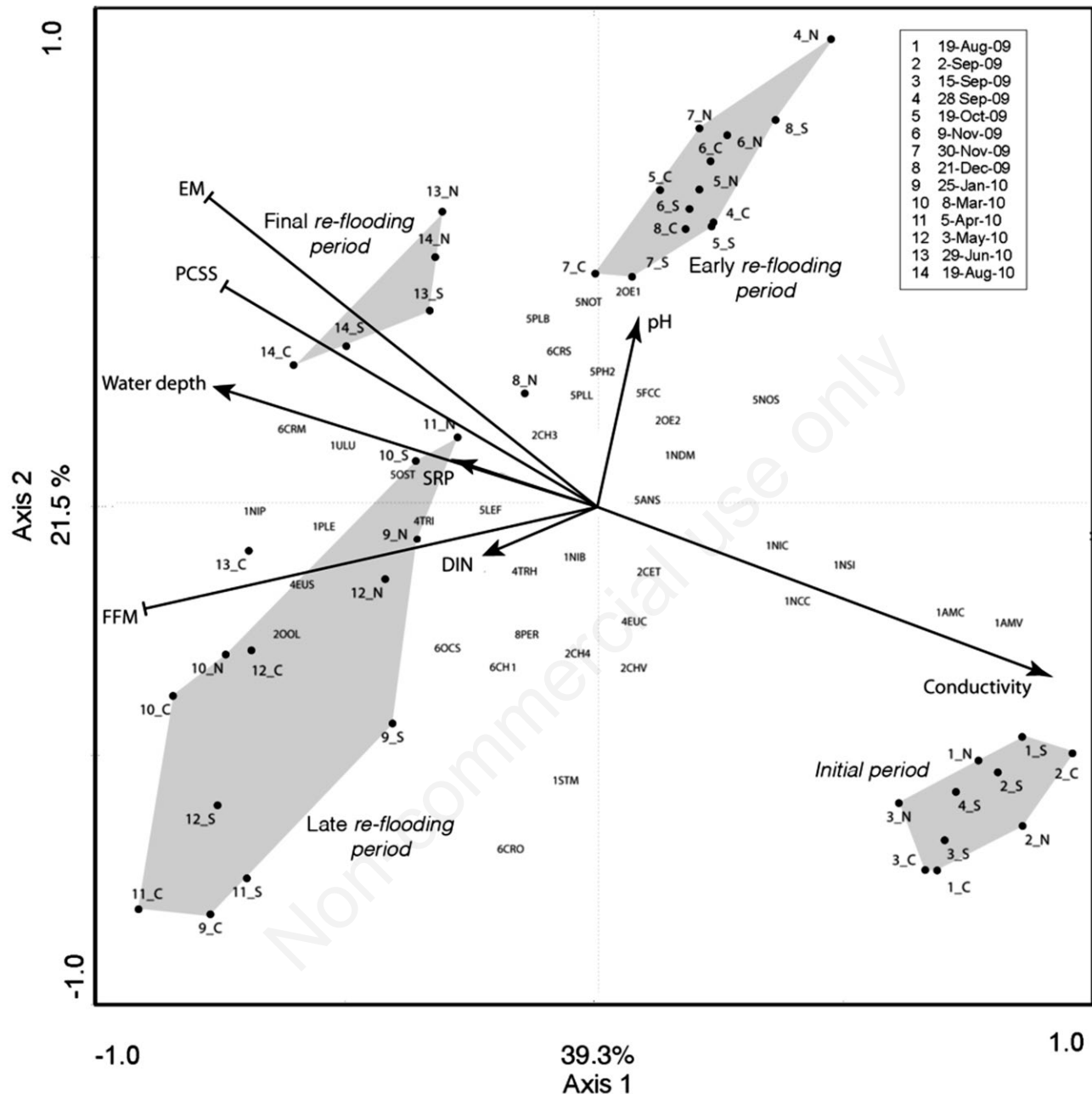
Niño and La Niña reflect the impact that these extreme hydrometeorological events may pose on wetlands.

During the studied annual cycle, both water level and macrophyte surface coverage fluctuated. Aquatic macrophytes play an important role in structuring communities in aquatic environments as they provide physical structure and increased habitat heterogeneity and complexity (Thomaz and Ribeiro da Cunha, 2010). Presence of macrophytes affects directly and indirectly the phytoplankton community by reducing water column mixing, limiting nutrient concentrations due to competition, reducing sediment resuspension and oxygenation of the water (Scheffer *et al.* 1993, 2003). In this respect, in our study the highest suspended solid concentrations in the water column coincided with the absence of vegetation. A profuse development of floating plants creates a stressful environment for phytoplankton, due to the low light intensity that impairs photosynthesis and growth, generating an imbalance between production and respiration (O'Farrell *et al.*, 2009). As suggested for other warm nutrient-rich wetlands (Meerhoff *et al.*, 2003; de Tezanos Pinto *et al.*, 2007), reduced nutrient concentrations (mainly DIN in our study) and anoxic environments underneath the free-floating macrophyte cover may explain the low phytoplankton biovolume registered. Phytoplankton Chl *a* concentration was typical of eutrophic and hypertrophic systems. As observed in other temporary water bodies (Cunha-Pereira *et al.*, 2010), peaks of Chl *a* occurred at varying times during the wet season. High values of Chl *a* were found at the beginning stages of flooding. The high values of this pigment measured in summer 2010 (second peak) can be explained by a concentration effect of algae, as a result of the reduction in the water level of Los Coipos Lake. This pattern is repeatedly encountered in

**Tab. 3.** Dominant phytoplankton species in terms of biovolume recorded in Los Coipos Lake and their functional group classification, throughout the study period (August 2009-August 2010).

	Site S	Site C	Site N
Initial period			
Species	<i>Euglena caudata</i> , <i>Cryptomonas cf. obovata</i> <i>Navicula cryptocephala var. veneta</i>	<i>Peridinium</i> sp.	<i>Stephanocyclus meneghiniana</i> , <i>Chlamydomonas</i> spp.
FG	W1, Y, T <sub>B</sub>	L <sub>O</sub>	C, X1
Early re-flooding period			
Species	<i>Trachelomonas hispida</i> , <i>Cryptomonas marsonii</i> , <i>Nitzschia palea</i>	<i>Cryptomonas cf. obovata</i> , <i>Euglena</i> sp., <i>Oedogonium</i> sp.	<i>Microcystis aeruginosa</i> , <i>Microcystis</i> sp., <i>Stephanocyclus meneghiniana</i>
FG	W2, Y, D	Y, W1, T	M, L <sub>M</sub> , C
Late re-flooding period			
Species	<i>Cryptomonas cf. obovata</i> , <i>Euglena</i> sp., <i>Stephanocyclus meneghiniana</i>	<i>Cryptomonas cf. obovata</i> , <i>Euglena</i> sp., <i>Mougeotia</i> sp.	<i>Cryptomonas</i> sp., <i>Microcystis smithii</i>
FG	Y, W1, C,	Y, W1, T	Y, L <sub>M</sub>

Site S, site South; Site C, site Central; Site N, site North; FG, functional groups proposed by Reynolds *et al.* (2002) and reviewed by Padisák *et al.* (2009).



**Fig. 6.** RDA ordination plot of samples based on their phytoplankton attributes. Asterisks indicate significant environmental variables. Numbers (1-14) indicate sampling dates. S, site South; C, site Central; N, site North, DIN, dissolved inorganic nitrogen; SRP, soluble reactive phosphorus; PCSS, percentage macrophyte-cover of site surface; EM, emergent macrophytes; FFM, free-floating macrophytes; 1AMC, *Amphora coffeaeformis*; 1AMV, *Amphora veneta*; 1NCC, *Navicula cryptocephala* var. *cryptocephala*; 1NIB, *Nitzschia brevissima*; 1NIC, *Nitzschia capitellata*; 1NDM, *Nitzschia dissipata* var. *media*; 1NIP, *Nitzschia palea*; 1NSI, *Nitzschia sigma*; 1PLE, *Pleurosigma* cf. *elongatum*; 1STM, *Stephanocyclus meneghiniana*; 1ULU, *Ulnaria ulna*; 2CH3, *Chlamydomonas* sp.3; 2CH4, *Chlamydomonas* sp.4; 2CET, *Chlamydomonas* sp.5; 2CHV, *Chlorella vulgaris*; 2OE1, *Oedogonium* sp.1; 2OE2, *Oedogonium* sp.2; 2OOL, *Oocystis lacustris*; 2PAM, *Pandorina morum*; 4EUC, *Euglena caudata*; 4EUS, *Euglena* sp.; 4TRH, *Trachelomonas hispida*; 4TRI, *Trachelomonas intermedia*; 5ANS, *Anabaena* sp.; 5LEF, *Leptolyngbya fragilis*; 5MIC, *Microcystis* sp.; 5MIA, *Microcystis aeruginosa*; 5MIS, *Microcystis smithii*; 5NOS, *Nodularia spumigena*; 5NOT, *Nostoc* sp.; 5OST, *Oscillatoria tenuis*; 5PH2, *Phormidium* sp.2; 5PLB, *Planktolyngbya brevicellularis*; 5PLL, *Planktolyngbya limnetica*; 5FCC, filamentous *Cyanophyceae*; 6CRM, *Cryptomonas marsonii*; 6CRO, *Cryptomonas* cf. *obovata*; 6CRS, *Cryptomonas* sp.; 6CH1, *Chrysophyceae* sp.1; 6OCS, *Ochromonas* cf. *viridis*; 8PER, *Peridinium* sp.

other Pampean shallow lakes close to being dry (Sosnovsky and Quiros, 2006), as well as in wetlands from other latitudes (Naselli-Flores and Barone, 2012). Small (<2  $\mu\text{m}$ ) fast-growing picophytoplankton is generally well represented in the phytoplankton community in shallow lakes of differing latitudes (Bell and Kalff, 2001; Callieri, 2007) and our results agree with this. Picocyanobacteria were one-fold greater than picoeukaryotic algae, which is a common finding in aquatic ecosystems worldwide (Callieri, 2007). Picoeukaryotic algae contribution is known to increase with decreasing light availability (Søndergaard, 1991; Callieri, 2007; Izaguirre *et al.*, 2010; Silviso *et al.*, 2011). In line with these findings, with the augmentation of *Azolla filiculoides* development in Los Coipos shallow lake, and concomitant reduction in light availability beneath the free-floating macrophyte cover, an increase in picoeukaryotic algae was also observed. Marked decreases in light penetration under profuse *Azolla* sp. coverage was measured in a nearby wetland (Reserva Natural Otamendi, Buenos Aires, Argentina) revealing that up 98% of total incident light was reduced under this situation (de Tezanos Pinto *et al.*, 2007).

As for the >2  $\mu\text{m}$  phytoplankton, statistical analyses evidenced the importance of environmental factors in shaping its assemblage during the *initial* and *re-flooding periods* of our study. In particular, changes in the depth of the water column, conductivity and development of an emergent and free-floating macrophyte cover affected the phytoplankton composition in Los Coipos Lake the most. Reynolds (2006) stressed that in truly novel planktonic habitats, the first colonists have to arrive *de novo*, and species that become initially abundant either *arrive* in strength or grow rapidly, or do both. Some unicellular organisms that have high growth rates are recognized as pioneer species. In this respect, following the functional group (FG) classification (after Reynolds *et al.*, 2002 and reviewed by Padisák *et al.*, 2009) we encountered small *Chlamydomonas* spp. (X1 FG, Chlorophyceae) and *Ochromonas* cf. *viridis* (X2 FG, Chrysophyceae) that achieved relatively high abundances in developing pools in the *initial period* of our study. It is worth mentioning that *Ochromonas* cf. *viridis* dominance was not expressed in terms of biovolume due to their relatively small size. These (X1 and X2 FG) are typical small fast-growing algae. We also found high relative proportions of euglenoids; it is recognized that the drying phase of temporary water bodies is characterized by a very diverse assemblage mainly formed by euglenoids (W1 and W2 FG) often accompanied by dinoflagellates ( $L_0$  FG) because they generally persist to the point of complete desiccation of the pond (Naselli-Flores and Barone, 2012). The dominance of benthic Bacillariophyceae (C and  $T_B$  FG) during the *initial period* was favored by the absence of macrophytes as it is recognized that heavy diatoms rely on tur-

bulence to counteract sedimentation (Padisák *et al.*, 2003; Allende *et al.*, 2009; Stević *et al.*, 2013).

In the *re-flooding period*, the new conditions triggered by the insipient formation of the continuous shallow lake allowed the further colonization and growth of planktonic algae as reflected by the replacement of benthic algae (mostly dominant in *initial period*) by pelagic ones. The dominance of Cryptophyceae on different sampling dates, can be explained by the fact that *Cryptomonas marsonii* and *C. cf. obovata* (Y FG), are typical mixotrophic flagellated organisms that can withstand occasional nitrogen and light limitation, as they can ingest bacteria (Tranvik *et al.*, 1989; Jones, 2000; Urabe *et al.*, 2000; Sinistro *et al.*, 2007; Unrein *et al.*, 2007; Allende *et al.*, 2009). This fact, and the capacity to migrate vertically in order to reach optimum light condition, allowed them to remain in the water column during the whole phytoplankton succession, and to reach a greater contribution when macrophyte cover was more profuse and/or DIN concentration was low in the late *re-flooding period*. Dominance of Cyanobacteria on certain occasions was related to the establishment of macrophytes, which results in low light conditions and enhanced physical stability of the water column. In this sense, Olding *et al.* (2000) stated that some cyanobacteria species are sensitive to vertical mixing. Also, it is known that some species of this group are able to grow at low light intensities and warm temperatures (Chorus and Bartram, 1999; Carey *et al.*, 2012). In particular, *Planktolychnya limnetica* (S1 FG) is tolerant of deficient light conditions (Reynolds *et al.*, 2002) and *Microcystis* (M or  $L_M$  FG) is a fast migrating genus that may benefit in stratified conditions by gaining a competitive advantage over other non-migrating or slow-migrating phytoplankton (Carey *et al.*, 2012). The commonest species of Chlorophyceae were nanoflagellated *Chlamydomonas* spp. and *Chlorella vulgaris* (both X1), which are both typically small pioneer organisms and were present throughout the study. These genera are amongst the major airborne algae (Happpy-Woods, 1988, and references therein; Genitsaris *et al.*, 2011) and they can rapidly colonize standing waters if the nutrient conditions are appropriate. However, due to their small size their contribution to the total biovolume was not significant. Different authors have suggested that different strategies adopted by phytoplankton are related to morphological (size, shape, motility) and physiological (nutritional requirements) differences (Reynolds *et al.*, 2002; Kamenir *et al.*, 2004; Carey *et al.*, 2012). In this sense under the extreme scenario encountered at the end of our study, which included reduced light availability and vertical mixing, species belonging to functional groups that have adaptations to withstand these harsh conditions, dominated the phytoplankton assemblage. Among them, typical flagellated mixotrophic algae - *Euglena caudata* (W1 FG),

*Ochromonas* sp. (X2 FG) and *Cryptomonas* cf. *obovata*, *Cryptomonas* sp. and *Cryptomonas marsonii* (all Y FG) - and Cyanobacteria - *Planktolyngbya limnetica* (S1 FG) and *Microcystis smithii* ( $L_M$  FG) - that are tolerant to low light conditions and are able to regulate buoyancy. As macrophytes grew during the *re-flooding period* epiphytic algae were also encountered in the water column. Detachment of these algae from their substrata as a result of wind action in low water periods affects species richness in the water column. Thus, during early *re-flooding period*, the increase in water depth and greater availability of substrata for epiphytes contributed to the augmenting of phytoplankton species richness. When conditions were more homogeneous and the whole shallow lake was covered by free-floating plants in the late *re-flooding period*, the conditions provoked a reduction in species richness as suggested for other freshwater ecosystems under similar conditions (Scheffer *et al.*, 2006). Values of the SIMI index revealed that initially the three sites (pools) strongly differed in their phytoplankton species structure showing a relatively high spatial heterogeneity, probably due to the fact that different resting stages of colonists were able to develop in the nascent community of the new planktic habitat in Los Coipos Lake. However, the relatively high SIMI index observed among sites at the end of the study evidenced that the structure of the phytoplankton assemblage was forced not only by the water level increase, but also by the strong effect imposed by the complete free-floating macrophyte cover.

As stated above, under the conceptual model of Goldsborough and Robinson (1996) for algae in wetlands, we recognize that during the drought/flood period studied, Los Coipos Lake was initially in a *dry state*. In our study, after the re-flooding a shallow lake was formed that ended-up being completely covered not by metaphyton as expected for the *sheltered state*, but rather, by emergent and free-floating plants. In this regards, the phytoplankton community was exposed to some of the environmental conditions prevalent under the *sheltered state*, such as presence of aquatic macrophytes, reduced light penetration, medium water level and stability of the water column. However, it must be pointed out that in this wetland it was provoked by the presence of aquatic plants and not metaphytic algae.

In conclusion, our results provide evidence that changes in water level and the complete surface coverage by free-floating macrophytes were the main determinants of the structure and dynamics of the phytoplankton assemblage. In particular, the functional group approach revealed that the phytoplankton assemblage responded to rapid water level shifts (mainly represented by small fast-growing X1 and X2) and to the changes imposed in the water conditions as a result of the presence of a total free-floating macrophyte cover ( $W_1$ , T y  $L_M$ ).

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