### Phytoplankton community as monitoring tool in the terminal stretch of a micro-tidal estuary facing the Tyrrhenian Sea

Francesco Bolinesi<sup>1</sup>, Emanuela Serino<sup>1</sup>, Angelo Carotenuto<sup>1</sup>, Silvia Fanina<sup>1</sup> and Olga Mangoni<sup>1,2</sup>

<sup>1</sup>Department of Biology, University of Naples Federico II, Complesso Universitario di Monte Sant'Angelo, Naples; <sup>2</sup>NBFC, National Biodiversity Future Center, Palermo, Italy

#### ABSTRACT

Mediterranean coastal systems, particularly those characterized by river mouths, have played an important role in the history of human civilization, thanks to unique species assemblages, a high biodiversity, and complex ecological interactions. The overlapping of multiple cross-scale effects, triggered by the ongoing climate change, makes these systems particularly important sites for ecological studies. Here, we have investigated the phytoplankton community structure in the terminal stretch of the Sele River (Tyrrhenian Sea) under two different hydrological regimes: i) with river mouth occluded by sediments preventing water exchange with the sea; ii) with river mouth completely opened, which allowed the maximum water mass exchanges between the river and the sea. The strong variations in physical-chemical

Corresponding author: Olga Mangoni, Department of Biology, University of Naples Federico II, Complesso Universitario di Monte Sant'Angelo, via Cinthia 21, 80126, Naples, Italy. E-mail: olga.mangoni@unina.it

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This work is licensed under a Creative Commons Attribution-NonCommercial 4.0 International License (CC BY-NC 4.0). properties of the water column, the point-distribution of total phytoplankton biomass, and the dominance of cryptophytes and cyanophytes under occluded and opened river mouth, respectively, indicate the presence of strong allogenic pressures acting on the system. Our data suggest that the monitoring of phytoplankton communities in the terminal stretch of micro-tidal estuaries, could be a useful tool for studying the extent of climate change underway in Mediterranean coastal marine areas.

### Introduction

Mediterranean micro-tidal environments are transitional systems characterized by a tidal range <2m, where salt-wedge intrusion along the final stretch of rivers plays an important role in shaping biological communities (Boothroyd, 1978; Davies, 1964; Kiriaki *et al.*, 2010; van der Tuin, 1991). The extent of this process depends on several geomorphological and physical factors, such as, for example, the river mouth configuration or man-made barrier-like dams (Haralambidou *et al.*, 2010; Lichter *et al.*, 2011).

From a geomorphological point of view, a fennel-shaped river mouth can be defined as an estuary, in which the mixing between seawater and freshwater generates strong salinity gradients and the presence of a typical brackish water layer (Perillo, 1995; Poulos et al., 1993; Valle-Levinson, 2010). These systems are particularly vulnerable to climate change, since strong seasonality and variability in precipitation produce large inter-annual fluctuations in river flow and water mass exchange with the sea (Gasith and Resh, 1999; Kennish, 2002, 2021; Newman et al., 2005; Ustin et al., 2014; Verri et al., 2018;). In the last decade, in fact, a reduction of net rainfall of ~33% of the annual mean was reported for southern Italian regions, mostly due to a reduced contribution of winter rainfall (Caloiero et al., 2021; Polemio and Casarano 2004, 2008). The effects of the drought period and human interventions like land reclamation, construction of river embankments, and water extraction activities significantly contributed to altering the ecological profiles of these areas (Dugdale et al., 2012; Hearn and Robson, 2001; Kennish, 2021; Newman et al., 2005; Richard et al., 2018;). The combined effect of saltwater intrusion and river discharge produces significant changes in nutrient loads, since the extent to which nutrients are retained in rivers and then made available during the biomass development



seasons depends on the water retention time (Lancelot and Muylaert, 2011; Lichter et al., 2011; Newman et al., 2005; Pinckney et al., 1999;). Many authors state that changes in river water regime from lotic to lentic are accompanied by an increase in allogenic phytoplankton biomass (Lampert and Sommer, 1997; Reynolds, 1984, 1999, 2003; Reynolds and Descy, 1996; Waylett et al., 2013) generating blooms under optimal environmental conditions dominated by species able to grow at intermediate salinity as well as tolerant to large salinity fluctuations in brackish waters of terminal stretches (Jackson et al., 1987; Muylaert et al., 2009; Roubeix and Lancelot, 2008). Being passively transported by currents, phytoplankton at the source of the river would require a very rapid replication time so that they can generate blooms. This occurs when net specific growth rates of species exceed the residence time of the water (Lucas et al., 2009; Reynolds, 1984). Hence, phytoplankton represents a significant component of primary production only in slow-moving rivers in lowland regions. The responses of the phytoplankton community to variations in water flow rate, nutrient loads, light availability, and temperature are species-specific and highly dependent on the whole environmental context, differing from time to time within each segment of an estuary (Artigas et al., 2014). Several studies report that diatoms are the dominant phytoplankton group in rivers, with changes in size spectra composition and species associated with changes in nutrient concentration, turbidity, and flushing rate (Genkal 1997; Reylonds et al., 1994b). For example, Schuchardt and Schirmer, (1991) and Muylaert et al., (2000) report the dominance of diatoms in turbid estuaries throughout the year under high nutrient availability. Marshall et al., (2006) report the dominance of diatoms in spring in estuaries where turbidity is low, when light levels are relatively low and stratification does not occur. Among other groups, dinoflagellates and cyanobacteria are typically more abundant in summer during periods of low water discharge, long water residence time, minimal flushing rates, and low turbidity levels (Lancelot and Muylaert, 2011). Tavernini et al., (2011) report the dominance of diatoms, chlorophytes, and cyanophytes in the terminal stretch of the Po River, with changes in seasonal species assemblages showing an increase of cryptophytes in late July and of cyanophytes at high water discharge rates, from the end of April to mid-June. Within the problematic framework described above, we have analyzed the phytoplankton community structure through the detection of diagnostic pigments (Ansotegui et al., 2001; Mangoni et al., 2017) in the final stretch of the Sele River (Tyrrhenian Sea) under two distinct hydrological regimes: during a severe drought period in July 2017, in which, for the first time in living memory, the mouth of the river was completely occluded by sediments, and in May 2021, in presence of a high water flow rate and a completely open river mouth. Our aim was to understand to what extent the strong environmental variations influenced the structure of the phytoplankton community in the terminal stretch of one of the most important river Southern Italy.

#### **Materials and Methods**

#### Study area

The Sele River is the second largest river in southern Italy in terms of average water volume after the Volturno. It is located within a wide alluvial coastal plain, with a drainage basin of 3235 km<sup>2</sup> and a solid flow of 500,000 m<sup>3</sup> y<sup>-1</sup> (Cocco et al., 1989). 64 km long and tributary of the Tyrrhenian Sea (Di Paola et al., 2014), it has the highest mean annual flow discharge among the rivers of Southern Italy (*i.e.*, ~69.4 m<sup>3</sup> s<sup>-1</sup>) (Magliulo et al., 2021). It has been an important site for ancient populations, with the first archaeological settlements dating back to the second millennium B.C. (Ferrara et al., 2010). The climate of its basis is characterized by prolonged warm and dry summers and wet and mild winters, with mean annual precipitation between 700 to 2000 mm, (average of 1180 mm) showing a marked spatial variation (Diodato et al., 2011). Over the past 150 years, the Sele estuary coastline has been affected by erosion. From 1870 to 1984, the coastline gradually receded, with the highest erosion rates occurring in the main areas around the river mouth. In the last 150 years, the Sele River and its catchment area have also been affected by increasing human activities that exerted a significant influence on the evolution of the area. The construction of an artificial drainage channel (Canale di Bonifacimento) (Alberico et al., 2012a, b), with a dense net of artificial drainage channels and the presence of the Dam of Persano strongly enhanced coastal erosion and altered the river environmental conditions (Alberico et al., 2012a, 2012b; Arienzo et al., 2020; Pappone et al., 2011;). The Dam of Persano, located 16.2 km from the river mouth, built between 1929 and 1932, created a basin of 1.5 million m<sup>3</sup>, which affected the downstream sediment deposition causing the consequent retreat of the coast (D'Acunzi et al., 2008; Freeman et al., 2007; Magdaleno et al., 2018). In most recent years, the significant geomorphological alterations of the Sele estuary and the presence of pollutants and conspicuous loads of organic matter deriving from anthropogenic activities have been documented, along with a pronounced accumulation of allogenic substances under low flow conditions caused by rainfall deficit (Arienzo et al., 2020; Benassai et al., 2015; De Rosa et al., 2022; Montuori et al., 2022;).

#### Sampling and environmental parameters

Water sampling was carried out in the terminal stretch of the Sele River, from the mouth up to 3.5 Km upstream, under two distinct flow-rate conditions (Figure 1a-c). The first sampling was carried out in July 2017 (Arienzo *et al.*, 2020) with the river occluded by sediments, a condition that persisted for 1 week, and the second sampling was carried out in May 2021, with the river's mouth completely opened. Stations were located taking into consideration the presence of drainage channels flowing into the river (Table 1).

At each station, 5 L of water were collected with a Niskin bottle and successively subsampled for the analyses of biological and chemical parameters, as described below.

For the determination of total phytoplankton biomass (Chl a), 50-500 mL of water (depending on the presence of particles in the samples) were drawn from the Niskin bottle and filtered onto GF/F Whatman glass-fiber filters (25-mm diameter) immediately cryopreserved in liquid nitrogen until the analyses in the laboratory.

At each station, water temperature and salinity were determined using Idromar XMAR212 (2017) and Sbe 19 plus (2021) CTD probes interfaced with a GPS (Garmin Map 78S, Garmin, Olathe, USA).

# Total phytoplankton biomass and chemo-functional groups

For the determination of total phytoplankton biomass (Chl a), 50-500 ml of water (depending on the presence of particles in the samples) were drawn from the Niskin and filtered on GF/F Whatman filters (25-mm diameter) immediately cryopreserved in liquid nitrogen until the analyses in laboratory (Holm-Hansen *et al.*, 1965; Lorenzen and Jeffrey, 1980; Yentsch and Menzel, 1963). Chlorophyll-a and phaeopigments were determined, after extraction with 90% acetone, with a spectrofluorometer (Mod.RF–6000; Shimadzu Corporation, Kyoto, Japan), with calibration curves checked daily with fresh Chl a standard solutions (Sigma-Aldrich, St. Louis, USA) (Holm-Hansen *et al.*, 1965). The pheao:chl a ratio was used as a proxy of grazing activity on phy-

 Table 1. Sampling station with coordinates and bottom depth.

Station	Latitude N	Longitude E	Depth (m)
1	40.4814990	14.9448934	3.037
2	40.4812454	14.9488489	6.017
3	40.4810787	14.9506658	3.819
4	40.4812777	14.9538731	3.221
5	40.4829086	14.9582883	3.333
6	40.4850757	14.9623601	3.479
7	40.4877350	14.9655997	3.157
8	40.4902284	14.9688533	3.306
9	40.4920247	14.9718688	3.414
10	40.4962250	14.9724679	2.732
11	40.4997458	14.9718300	3.559





Figure 1. Final stretch of the Sele river with sampled stations (a). Mouth of the river in July 2017 (b) and May 2021 (c).

toplankton cells (Shuman and Lorenzen, 1975). For the determination of the pigmentary spectra, 1.5 L of water was filtered on GF/F Whatman filters (47-mm dimeter) and the filters were cryopreserved as for Chl a. Frozen filters were homogenized and resuspended in 100% methanol and analyzed by High-Performance Liquid Chromatography (HPLC) (1100 Series, Hewlett Packard, Palo Alto, USA) in a reverse phase (C8 column 3 µm Hyperloop MOS) (Vidussi et al., 1996). The use of HPLC in the study of phytoplankton communities has been largely demonstrated to be a useful tool for the estimation of phytoplankton community composition through the analyses of photosynthetic pigments (Jeffery and Vesk, 1997; Wright et al., 1996). The method is based on the analysis of accessory pigments, in addition to Chlorophyll-a (Chla) or the modified divinyl-Chl-a found in all phytoplankton species, and on the evidence that some of these accessory pigments are taxon-specific (Brunet and Mangoni, 2010; Millie et al., 1997; Wright and Jeffrey, 2006). This technique allows to detect and identify microscopically overlooked or undetermined ultraphytoplankton species (Ansotegui et al., 2003; Antajan et al., 2004; Garibotti et al., 2003; Saggiomo et al., 2023), providing reproducible results. For the determination of chlorophylls and carotenoids, a spectrophotometer with a diode array detector was set at 440 nm, making it possible to determine the absorption spectrum of the 350-750 nm interval for each peak to check the purity of single pigments. The column was calibrated using different pigment standards (chlorophyll a, chlorophyll b, chlorophyll  $c_1+c_2$ , chlorophyll c<sub>3</sub>, MgDVP, alloxanthin, diatoxanthin, prasinoxanthin, 19'-butanoyloxyfucoxanthin, fucoxanthin, 19'-hexanoyloxyfucoxanthin, peridinin, zeaxanthin, neoxanthin, violaxanthin, bcarotene). Standards were provided by the International Agency for 14 C Determination, VKI Water Quality Institute, Copenhagen, Denmark. Quantification was based on the absorbance at 440 nm and the factor response value (peak area/pigment concentration) for each pigment, as described by Mantoura and Repeta (1997). The contribution of main phytoplankton groups to the total Chl a was estimated by CHEMTAX 1.95 software as indicated by Latasa (2007), using an iterative process to find the optimal pigment: Chl a ratios. The chemotaxonomic groups identified include cyanophytes (Cyano), chlorophytes (Chloro), prasinophytes (Prasino), euglenophytes (Eugleno), cryptophytes (Crypto), diatoms (Diato), pelagophytes (Pelago) haptophytes (Hapto), dinoflagellates (Dino), xantophytes (Xanto).

#### **Inorganic nutrient concentrations**

For the determination of inorganic nutrient concentrations  $(N-NO_3, P-PO_4)$ , 10 mL water were filtered on 0.2 µm cellulose acetate filters, and stored in HDP vials at -20C. The analyses in the laboratory were conducted using a discrete sampling analyzer (EasyChem Plus, Systea, Anagni, Italy) equipped with a UV-VIS spectrophotometer following the procedure described by Hansen and Grasshoff (1983) adapted to current instrumentation, using a standardized method to detect nitrate (EPA - Ref. National Environmental Methods Index 9171 Nitrate via V(III) reduction), and orthophosphate (ISO 15923 - Ref. International Standard Organization 15923-1 Water quality).

#### Statistical analyses

The spatial distribution of physical variables (temperature, salinity) was plotted using ODV 5.6.3 software.

To estimate the pairwise dissimilarity between the two contrasting water flow conditions in a low-dimensional space, a non-metric multidimensional scaling based on a distance matrix computed with the Bray-Curtis dissimilarity has been performed and represented using a bi-plot in which the correlation coefficients between environmental variable and the NMDS scores were presented as vectors from the origin based on algorithm proposed by Taguchi and Oono (2005). The significance of differences between the two putative conditions of river's water flow was tested by ANOSIM (Clarke, 1993). All statistical analyses were performed using the PAST 1.95 software.

#### **Results**

#### **Environmental parameters**

Values of salinity are reported in Figure 2. In 2017 the distribution of salinity showed the presence of a strong water stratification, with a net halocline between 1.5-2 m (station 2) and 2-2.5 m (station 9) whose depth increased moving toward the inner part of the river (Figure 2a). In general, values ranged between 39.8 and 0.7. At station 1, near the sandbar occluding the river mouth, salinity reached values up to 39 originating a 'bubble-like' front probably related to the seawater percolation or episodic waves intrusion from the adjacent sea. The isohalines between 5 and 8, which have been the subject of dispute and debate over time (Deaton and Greenberg, 1986; Khlebovich, 1968, 1969), took place between 1 m (station 2) and 1.8 m depth at station 9, with a slightly deepening moving from the mouth to the inner part of the river. Station 1 was the only one with salinity >8 in the entire water column. In 2021, with the river mouth completely opened, the entire sampling area was still characterized by a net water stratification, with a 5-8 salinity boundary cline placed between 1.5 and 2.5 m depth, deepening moving from the mouth (station 1) to the inner part of the river (station 11) (Figure 2b). In the entire area, salinity ranged between 36.2 and 0.01. All stations presented two distinct layers: a freshwater one, between 0 and 1.5 m, and a marine one, below 2.5 m. The only exception was station 10, where values reached maximum salinity of 10 on the sea bottom, and the freshwater laver reached its maximum thickness.

As far as the temperature is concerned, it showed clear differences between the two sampling periods (Figure 3). In July 2017 water temperature exceeded 28.5 °C in the first meter of the water column, generating a marked thermocline at  $\sim 1.3$  m depth in the overall sampling area (Figure 3a). The bottom layer was characterized by the presence of the coldest water reaching the minimum of 24.59 °C at station 2. The surface layer showed a slight increasing trend of temperature moving from the mouth to station 11, with values ranging between 28.5°C and 31.34°C. In May 2021, the water column appeared to be rather homeothermic (Figure 3b). In general, values decreased moving from the mouth to the station 11, where temperature reached the minimum of 19.3 °C. A weak hermos-stratification was present in the first 30 cm between stations 1 and 5, where the water retention time reached its maximum. Nevertheless, a slightly increased temperature in the bottom water layer was observed in vertical profiles between stations 5 and 9, in accordance with the increase of salinity linked to seawater inflow.

The concentration of N-NO3 and P-PO4 in July 2017 led to

a mean N:P ratio of 12.12 in the entire area, showing a pointdistribution depending on the influence of drainage channels (Arienzo *et al.*, 2020). In particular, N-NO<sub>3</sub> ranged between 12.85  $\mu$ mol L<sup>-1</sup> (station 5) to 23.80 (station 10) with a mean concentration of 18.2  $\mu$ mol L<sup>-1</sup>; P-PO<sub>4</sub> ranged between 0.81  $\mu$ mol L<sup>-1</sup> (station 3) and 8.39  $\mu$ mol L<sup>-1</sup> (station 10) with a mean concentration of 2.84  $\mu$ mol L<sup>-1</sup> (Figure 4a).

In May 2021, values of N-NO<sub>3</sub> showed a mean concentration of 15.76  $\mu$ mol L<sup>-1</sup> ranging between 4.57  $\mu$ mol L<sup>-1</sup> (station 2) and 23.51  $\mu$ mol L<sup>-1</sup> (station 9), and displaying a point-shaped distribution, as observed in the first sampling (Figure 4b, c). P-PO<sub>4</sub> were characterized by very low concentrations, with a mean of 0.2  $\mu$ mol L<sup>-1</sup> and values ranging between 0.08  $\mu$ mol L<sup>-1</sup> (station 2) and 0.39  $\mu$ mol L<sup>-1</sup> (station 6). These aspects led to a mean N:P ratio of 83, a value far from what was observed in 2017 and from the Redfield one. In particular, the N:P ratio showed values ranging between 148 (station 3) and 30 (station 5), with the highest values in the surface layer.

# Phytoplankton biomass and chemo-functional groups

The distribution of total phytoplankton biomass (Chl a) in July 2017 showed the presence of eutrophic condition in the





overall sampling area (Figure 5a). Values ranged between a minimum of 15.05  $\mu$ g L<sup>-1</sup> at station 9, and a maximum of 95.73  $\mu$ g L<sup>-1</sup> at station 10, highlighting a strong variability in the area, especially near drainage channels. This aspect was also emphasized by the fluctuation of grazing index values (Phaeo:Chl a), which showed a mean of 0.59 ranging between 0.20 (station 11) and 2.54 (station 9). The overall picture that emerged from Chl a concentration indicated the presence of an increasing trend moving from the mouth to the inner part of the river. In May 2021 (Figure 5b, c), values of Chl a were at least one order of magnitude lower compared to what was reported before, with values usually observed in oligotrophic/mesotrophic systems. The mean value for the entire area was 1.40  $\mu$ g L<sup>-1</sup>, with concentrations ranging between 0.55  $\mu$ g L<sup>-1</sup>(station 10) and 3.11  $\mu$ g L<sup>-1</sup>(station 11). The grazing index ratios were very high at all stations, with a mean of 1.21 and values ranging between 0.62 (station 2) and 1.87 (station 10).

As far as the pigment spectra composition (expressed as pigment:Chl a ratios) is concerned, we observed high variability in both sampling periods (*Table S1*). In July 2017, fucoxanthin was the dominant pigment, with a mean ratio of 0.67 and values ranging between 0.43 (station 5) and 0.97 (station 1). Alloxan-



Figure 3. Distribution of temperature along the water column of the entire sampled area in July 2017 (a) and May 2021 (b). Y axis: depth (m); X axis: station distance; Z colored axis: temperature ( $^{\circ}$ C).

thin showed the highest ratio at stations 5 (0.64) and 10 (0.64)and, together with Chl c2 and Diadinoxanthin was among the most represented ones. Chl c3, peridinin, phaeophytin, and 19'HF were the only pigments completely absent in the sampling area. In May 2021, fucoxanthin and MgDVP were the most abundant pigments, with mean ratios of 0.49 and 0.41, respectively. aFucoxanthin ranged between 0.36 (station 2, bottom layer) and 0.56 (station 5, bottom layer); MgDVP between 0.29 (station 4) and 0.59 (station 8, bottom layer). The contribution of fucoxanthin and MgDVP to the total pigmentary spectra was higher at the bottom layer than 0 m. Neoxanthin was the only pigment to be absent at all stations. Peridinin and neoxanthin were completely absent only at surface layer, while other pigments showed weak and nonlinear changes as highlighted by colored bars. The pigmentary spectra composition descripted so far, determined a different chemotaxonomically composition of phytoplankton community in the two sampling periods (Figures 6a-c). In July 2017, cryptophytes strongly dominated the community in the entire area, with a mean of 58% and values ranging between 29% (station 2) and 79% (station 11). Diatoms and chlorophytes were the second most representative groups, with a mean of 16% and 11% respectively, and higher percentages at station 7. Dinoflagellates and haptophytes were completely absent, while xanthophytes were only reported with a weak percentage (2%) at station 11.

In May 2021 (Figure 7b-c), the community was dominated by cyanophytes and diatoms, with mean percentages of 28.5% and 27.3% respectively. Cyanophytes showed a strong gradient at station 4, ranging between 9% (surface layer) and 43% (bottom layer); diatoms ranged between 6% (station 3, bottom layer) and 47% (station 11). Chlorophytes and euglenophytes showed similar mean percentage (7%), with the first group that was more represented at the surface layer, and the second one more represented at deeper layer where reaches percentages up to 16% (station 11). Haptophytes was the only groups to be absent at surface layer and to be present exclusively at bottom of the station 4.



**Figure 4.** Box plots with nutrient concentrations of P-PO4 and N-NO3 in July 2017 (a); surface layer in May 2021 (b); bottom layer in 2021 (c).



**Figure 5.** Chart bar with chlorophyll a concentration (Chl a  $\mu$ g/L) and grazing index (Phaeo:Chl a) at each station (X axis) in July 2017 (a); surface layer May 2021(b); bottom layer in May 2021(c).



#### Discussion

Mediterranean microtidal estuaries have played an important role in human civilization (Haidvogl 2018). The present study aims at providing the first insights into changes in phytoplankton community structure in the terminal stretch of the Sele River, a microtidal estuary subjected to multiple anthropogenic pressures, as the result of contrasting conditions of the river's water flow.

The reclamation works carried out in the region during the last century have profoundly altered its morphology and degree of interactions between the river, the surrounding lands, and the sea. The extent of these changes is reported today in several geomorphological studies, though with a considerable lack of information on the ecology of the area, especially regarding microalgal communities in the terminal stretch, where phytoplankton plays a pivotal role as primary producers in lentic regime waters. Our data represent the first attempt to describe the phytoplankton community structure in one of the most important hydrographic basins of southern Italy, under two distinct hydrological phases: with the mouth of the river completely occluded by sediments - for the first time in living memory–preventing the water exchange with the sea, and under a high hydrological regime with mouth completely opened.

Under two different hydrological conditions, the terminal stretch of the Sele River was characterized by the coexistence of two distinct water layers. A shallow one, showing typically freshwater salinity whose thickness tends to be higher in the inner part

**Figure 6.** Chemotaxonomically composition of phytoplankton community in July 2017 (a); surface layer in May 2021 (b), bottom layer in May 2022 (c).



**Figure 7.** Non-metric multidimensional scaling based on Bray-Curtis index. Environmental variables are shown in green; the correlation coefficients between each environmental variable and the NMDS scores are presented as vectors from the origin. Stations of July 2017 are reported as black circles; stations of May 2021 are reported as filled (bottom layer) and empty (surface) orange squares.

of the river, and a deeper one with salinity values that typically exceeded that of the adjacent seawater, especially in deepest ponds and during summer. These thickness changes were accompanied by changes of 5-8 isohaline depth, which was absent near the mouth occluded by sediments and highest at station 10 in the inner part of the river. Salinity variation represents a key factor for the biotic component of ecosystems and the presence of brackish organisms is a typical property of estuarine environments. Changes in salinity over river mouths depends on the tides, coastal flooding, and season of the year (Reynolds, 1984; 2006), so the formation of sandy bar occluding the mouth and preventing the water exchange with the sea, can have significant effects on the functioning and ecology of such a complex system, with multi-scale effects still unknown.

Since dramatic rainfall-decreasing trends were reported for the southern Italian Tyrrhenian coast, causing droughts during summer (Polemio and Casarano, 2004, 2008; Romano *et al.*, 2022), the results presented here could be the first sign of a new condition towards which this system will likely tend more frequently in the near future.

Contrarily to the salinity, the thermal profile of the water column showed marked differences between the two sampling periods in accordance with the season. In July 2017, a net thermocline was present at  $\sim 1.3$  m depth, with water temperature exceeding 28.5 °C in the first meter of the water column reaching colder values on the bottom (minim 24.59 °C, at station 2). In spring 2021 the water column was rather homeothermic, with lower values in the inner part of the river and relatively warmer water on the bottom linked to the seawater intrusion. A weak stratification was observed in the first 30 cm between stations 1 and 5, and despite the fact current meters data are not available, we hypothesize that this warmer layer reflects the hydrological dynamics of the area, where the increase in water retention time favors surface waters warming. It must be noted that stations were sampled ~10 minutes apart from each other, moving from the mouth to station 11, so sampling activities should not have influenced these data.

Many authors indicate that any potential alteration to river flows would increase the temporal and spatial variability of estuarine fronts, water stratification, and mixing, with potentially negative impacts associated with eutrophication (Reynolds, 1984; 2006). The impact of nutrient enrichment on rivers is often season specific and complicated by their dynamic nature (Newman et al., 2005). Differences in nutrient concentration observed in our study, with extremely large spatial and temporal variations in N:P ratios, suggest how complex and articulated are the dynamics of nutrient supply in this area, where the presence of drainage channels (in addition to the combined effects of water retention times) can strongly influence the chemical properties of the water column (Albanese et al., 2007; Arienzo et al., 2020; Montuori et al., 2022). Nitrates did not show drastic changes between the two sampling periods, as instead observed with the phosphate load, which led to higher N:P ratios in 2021, with values up to 148 at station 3, compared to those in 2017. The first biological effect of nutrient enrichment in pelagic environments is the growth of phytoplankton species, whose blooms extent and community structure strongly depend on local conditions, for example, the extent to which the nutrient load is retained by the receiving water (Newman et al., 2005; Reynolds, 2006). This is the reason why rivers are considered highly selective environments, where flow rate and turbidity, along with temperature, are the most critical factors limiting the development and structuring the phytoplankton community composition (Rojo et al., 1994; Reynolds, 1994, Tavernini et al., 2011). The higher the flow rate and turbidity, the lower the amount of phytoplankton can be expected: where the water retention time is longer than the generation time of the phytoplankton, large blooms can occur (Ibeling et al., 1998; Lampert and Sommer, 1997; Newman et al., 2005). Nevertheless, the extent to which nutrient loads trigger phytoplankton blooms in micro-and macrotidal estuaries is different, with small nutrient enrichment producing larger effects in the microtidal ones (Warwick et al., 2018). In general, allogenic factors (i.e. inoculum of phytoplankton species transported downstream, temperature, light regime, discharge rate, turbidity) are considered the main drivers of phytoplankton succession in lowland rivers (Reynolds, 1994, 2006), with autogenic ones becoming more relevant with increasing eutrophication levels (del Giorgio et al., 1991).

The distribution of total phytoplankton biomass (Chl a) in July 2017 points to a speckled distribution with an increasing trend of concentration moving toward the inner part of the river, where values are typical of eutrophic waters reaching the maximum of 97.73 µg L<sup>-1</sup>. The station with lowest concentrations of Chl a (15.0  $\mu$ g L<sup>-1</sup>) showed the highest phaeo:Chl a ratio (2.7), indicating the presence of large amounts of degraded biomass, most likely because of the adjacent drainage channel transport. On the contrary, in May 2021, values of Chl a were at least one order of magnitude lower, with a mean of 1.40 µg L<sup>-1</sup>, appearing more homogeneously distributed in the entire sampling area and being characterized by very high phaeo: Chl a ratios, the latter supporting the hypothesis that much of the phytoplanktonic biomass was photosynthetically inactive. The overall picture emerging from nMDS analyses clearly shows the differences between the two samplings periods, with data from July 2017 (black points) clustering in the left side of the plot, and those from May 2021 (orange squares) in the right part. Temperature, P-PO<sub>4</sub> and the N:P ratios are among the main environmental drivers shaping the phytoplankton community, with high Chl a concentration and cryptophytes characterizing the first sampling period, and high N:P ratios characterizing the second one. In May 2021 (orange squares), euglenophytes, cyanophytes, and xantophytes were the dominant groups, especially at the surface layer (filled orange squares). The correlation between salinity and depth highlights the role of seawater inflow in shaping the structure of the phytoplankton communities, with dinoflagellates, haptophytes, and pelagophytes dominating in the deepest and saltiest layers, and high loads of N-NO3 that were instead associated to the shallow layer, because of inputs from drainage channels. A high and positive R value (0.9777) in the ANOSIM analysis (p=0.0001) indicates the strong and significant dissimilarity between the phytoplankton communities' composition in the two sampling periods (Table 2). Altogether our data

 Table 2. Results of the ANOSIM (ANalyses Of SIMilarities)

 test carried out to ascertain differences between the two sampling periods, representing contrasting water flow conditions.

ANOSIM Test – Bray-Curtis				
Permutation N	9999			
Mean rank within	132			
Mean rank between	334.6			
R	0.9777			
p (same)	0.0001			

agree with the results reported by Tavernini *et al.*, (2011), who, in the Po River, observed the highest concentration of cryptophytes biovolume in late July, concomitantly with an higher contribution of cyanophytes between April and June: these patterns provide a further confirmation that the terminal stretch of the Sele River is exposed to strong allogenic pressures.

#### Conclusions

In addition to what discussed so far, we pinpoint here that the phytoplankton community represents a promising and effective monitoring tool for the study of estuarine environments, in which cross-scale interactions produce large fluctuations in physical, chemical, and biological properties. Nevertheless, the large variability of the phytoplankton community structure both in space and time, and the extremely ample variations in N:P ratios underline how much allogenic pressures can influence communities' functional traits in the terminal stretch of the Sele River and, supposedly, in other estuarine areas, with similar combinations of natural and anthropogenic pressure. The presence in both study periods of a saltier layer near the bottom highlights the role played by the seawater inflow in the ecology of the area, characterized by a strong summer thermoaline stratification, which exacerbates the difference between superficial and bottom water layers.

Although our results are not sufficient to mechanistically explain the functioning of such a complex ecosystem, the ample array of interactions among local climate, hydrodynamical and ecological variables that emerged from our study suggest that the Sele River estuary will most likely be increasingly exposed to the multiple stressors caused by the ongoing climate change and the increase of human uses of land and rivers. The presence of other estuarine environments in the Mediterranean Sea and the peculiar characteristics of the terminal stretches of rivers altogether suggest that more effort should be paid to investigate and monitor these delicate and vulnerable ecosystems. This appears to be urgently needed in order to identify new management measures of freshwater inputs in the Mediterranean Sea, one of the marine regions worldwide most prone to the consequences of climate change.

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Online supplementary material:

Table S1. Pigments: Chl a ratios at each station in July 2017 (a); surface layer May 2021(b); bottom layer in May 2021(c).