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**Mini lavori della Riunione scientifica del
Gruppo di Lavoro per l'Algologia**

(a cura R. Pistocchi)

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In copertina: aggregati di cellule di *Ostreopsis cf. ovata* (Dinophyceae) isolate nel Golfo di Trieste, con abbondante mucillagine formata da filamenti di tricocisti e polisaccaridi extracellulari (Golfo di Trieste)
foto di Giorgio Honsell

On the cover: aggregates of *Ostreopsis cf. ovata* (Dinophyceae) cells, isolated in the Gulf of Trieste, showing abundant mucilage composed of trichocyst filaments and extracellular polysaccharides (Gulf of Trieste)
photo by Giorgio Honsell

Factors affecting growth and toxin production in *Prorocentrum hoffmannianum* (Dinophyceae) in Florida Keys

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Prorocentrum hoffmannianum M.A.Faust is a benthic dinoflagellate that produces biotoxins which are causative of diarrhetic shellfish poisoning (DPS) in warm marine waters. The occurrence of this dinoflagellate was reported only in tropical areas (Belize, Florida, Mexico, Brazil, northern Australia and Hainan Island, China).

In this study, we investigated the effect temperature and nutrient depletion on *P. hoffmannianum* isolated in from field samples (Florida Keys, Florida, USA) in 2014.

Monoclonal cultures were prepared by isolating single cells by the capillary pipette method. Cultures were maintained at a temperature of 21 ± 0.1 or 26 ± 0.1 °C and a light intensity of $90\text{--}100\ \mu\text{mol m}^{-2}\ \text{s}^{-1}$ with light-dark cycle 12-12 h. For each temperature, *P. hoffmannianum* were grown in f/4 medium (as control), N-free and P-free medium in triplicate. Subsamples (2 ml) were sampled every 2 days for 30 days. Cell counts were made following the Utermöhl sedimentation method, through epifluorescence microscopy.

Both temperature and nutrient condition significantly affected growth rates and maximum yield of *P. hoffmannianum* showing the maximum values recorded in high temperature and replete medium. Production of okadaic acid was induced in all conditions (min 50 ± 46 pg/cell; max 643 ± 189 pg/cell), showing values up to 1 order of magnitude higher than that observed in other DSP species.

Toxin concentration (both on a per cell and a biovolume basis) resulted enhanced at 26 °C and in P depletion, corroborating the knowledge that toxin production is modulated by cell physiological conditions, which are affected by a wide spectrum of factors, including several environmental stressors such as nutrient availability.

Toxin fraction released in the growth medium was neglectable. No okadaic acid esters were detected in this strain of *P. hoffmannianum*.

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***Ostreopsis ovata* Fukuyo 1981 along the eastern Adriatic coast**

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Ostreopsis spp. are benthic dinoflagellates that are known to cause harmful algae blooms. They are known producers of a range of palytoxins and ovatoxins. During bloom events those toxins can be found in high concentrations in benthic biofilms, in the water column as well as in coastal aerosols. *Ostreopsis* spp. distribution and related harmful algae blooms are well documented along the western Adriatic coast and parts of the western Mediterranean. Little was known so far about its distribution along the eastern Adriatic coast which presents very different ecological conditions with respect to the western Adriatic coast. A high resolution sampling regime along the eastern coast of the northern Adriatic revealed a dense population structure and high abundances along that coast line. Samples from the middle and southern Adriatic confirmed the presence of *Ostreopsis ovata* along the entire eastern Adriatic coast. Our results demonstrate, that oligotrophic rocky shores can be a very suitable habitat for *O. ovata*. Massive bloom events with record abundances and toxin concentrations are reported. The population along the north-eastern Adriatic coast share one ITS1-5.8S-ITS2 genotype with most of the populations recorded around the Mediterranean.

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Hydrogel scaffolds from *Trichormus variabilis* (Nostocales, Cyanobacteria) exopolymers

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Cyanobacterial Extracellular Polymeric Substances (EPS) are highly complex and variable compounds exhibiting hydrophilic and anionic properties. EPS can comprise heteropolysaccharides, of six or more sugar units, proteins, nucleic acids and lipids. EPS highly retentive properties make them promising for producing hydrogel scaffolds for drug delivery and tissue engineering.

Released-EPS (R-EPS) were obtained by intensive indoor cultivation of a strain, VRUC 168, of the cyanobacterium *Trichormus variabilis* (Kützing ex Bornet & Flahault) Komárek & Anagnostidis, isolated from sediments of a Mediterranean coastal lagoon, in polyethylene bags (10L; BG11 medium; 25 °C; L-D 12-12h; irradiance 80 $\mu\text{mol photons/m}^2/\text{s}^1$). At stationary phase, day 20, the culture medium was collected and R-EPS concentrated by centrifugation and rotary evaporation. R-EPS were then precipitated in cold ethanol and the pellet freeze-dried. This material was characterized by RP-HPLC, spectroscopic analyses, cytochemical staining, BCA and phenol-sulphuric acid assays.

R-EPS were used to produce hybrid hydrogels (EPS-Hys) for detoxification enzyme- and stem cells- carrier systems. To this end, a specific protocol for the synthesis of the EPS-hydrogel through photopolymerization was developed. Hydrogel stability, resistance to hydrolysis and mechanical properties were also assessed. Thus, hybrid hydrogels were tested for enzymatic encapsulation using thiosulfate-cyanide sulfur transferase (TST) and 3D scaffolding for human cardiac mesenchymal stem cells (cMSCs).

Mass cultures of *T. variabilis* reached productivity of $0.039 \pm 0.001 \text{ g(DW)L}^{-1}\text{d}^{-1}$ and the R-EPS yield was of 2 gDW. Characterization of this EPS fraction revealed the presence of carbohydrates with sulfated and carboxylic groups. FT-IR spectrum showed peaks within 2900 and 3450 cm^{-1} , attributable to -CH and -OH group vibrations, and a peak around 1600 cm^{-1} attributable to -COOH group vibrations.

The fabricated hybrid hydrogels (EPS-Hys) resulted more stable and resistant to dehydration and spontaneous hydrolysis, at 37 °C, than PEGDa-Hys and rheological analysis of hybrid hydrogels evidenced that cyanobacterial EPS improved hydrogel mechanical performances.

Enzymatic encapsulation test showed that TST activity was preserved in the hydrogel at 37 °C. 3D scaffold application of the novel hybrid EPS-Hy showed effective adherence and growth of cMSCs cells, with an increase in cell viability after 3-day growth, suggesting a non-cytotoxic effect of the EPS-Hys on the stem cells. Results showed that *T. variabilis* S-EPS are a sustainable source of biomaterials adaptable to a large variety of biotechnological applications.

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Two new species of the genus *Ceramium* Roth (Ceramiaceae, Rhodophyta) from the Venice Lagoon (Italy)

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The cosmopolitan genus *Ceramium* Roth (Ceramiaceae), with about 213 currently accepted species (and infra-specific) names, is one of the largest groups in the Rhodophyta. This taxon is characterized by cylindrical or slightly compressed thalli, whose axial cells can be incompletely to completely covered by cortical cells, with alternate to pseudo-dichotomous branching, and straight to inrolled apices. It typically occurs in eulittoral or shallow subtidal habitats and its worldwide distribution is often connected with naval traffics. In fact, *Ceramium* thalli are often carried by hull fouling.

The nomenclature and the taxonomy of this genus are still in a state of chaos. Taxonomic problems are, indeed, tied to a high degree of variations in the morphological characters classically used in species recognition: presence of cortical spines, numbers of periaxial cells, developmental patterns of the corticating filaments, branching pattern, and tetrasporangial features. At the same time, the identification of new species is often complicated by their small sizes and epiphytic life style, that make them cryptic.

Culture studies, suggesting a strong influence of the environment on morphology, and the use of molecular tools have questioned the validity of morphological features to discriminate different *Ceramium* species. In particular, as the inventory of *Ceramium* species in some areas, has been only based on morphological observations, more and more authors have started using molecular markers (e.g. *rbcL* gene and *rbcL-rbcS* intergenic spacer, SSU rDNA, ITS2) to determine species diversity.

Here we characterize different isolates, sampled in transitional waters of the Venice Lagoon (Italy), in order to better investigate presence and biodiversity of *Ceramium* in these areas. Through accurate morphological observations and phylogenetic analyses, based on the plastid ribulose-1,5-bisphosphate carboxylase/oxygenase gene (*rbcL*) as molecular marker, we identify two distinct *Ceramium* taxonomic entities, never reported before in this environment according to the available checklists. The validity of the two entities as recognizable species is also discussed.

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Monitoraggio del fitoplancton lungo la costa laziale (Tirreno centrale) negli ultimi quindici anni

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Le attuali conoscenze della dinamica del fitoplancton marino nelle acque costiere del Lazio sono scarse. Poche sono infatti le pubblicazioni, risalenti ad oltre un decennio fa, incentrate sulla composizione della comunità fitoplanctonica e sulle sue variazioni spazio-temporali nelle acque superficiali, entro i 500 m dalla costa. I lavori nati dalla collaborazione tra l'Agenzia Regionale di Protezione dell'Ambiente, ARPA Lazio e il Laboratorio di Biologia delle Alghe dell'Università di Roma "Tor Vergata", hanno portato ad un primo aggiornamento dei popolamenti fitoplanctonici con l'identificazione di oltre 250 *taxa*, comprendenti specie tossiche, identificati con tecniche di microscopia ottica ed elettronica (Bianco et al 2006, Congestri et al. 2004, 2006). Il presente lavoro ha permesso di riprendere lo studio della componente fitoplanctonica del Lazio mediante la revisione e l'allineamento dei dati raccolti dal 2002 al 2016 nei programmi nazionali di monitoraggio, ai sensi della L. 979/82, del D.Lgs 152/2006 (direttiva quadro sulle acque 2000/60/CE) e del D.Lgs 116/2008 (direttiva sulle acque di balneazione 2006/7/CE), a cui sono stati aggiunti i dati prodotti nell'ambito della nuova attività di monitoraggio effettuata in ottemperanza al D.Lgs.190/2010 (direttiva sulla strategia marina 2008/56/CE). Tale programma, a tutt'oggi in corso, è stato avviato nel luglio 2015 e prevede un campionamento bimestrale lungo 4 transetti, ciascuno comprendente 3 stazioni di campionamento. L'aspetto innovativo di questo monitoraggio è la stima quali-quantitativa del fitoplancton anche in acque profonde in corrispondenza del DCM (Deep Chlorophyll Maximum) e più a largo, ovvero in stazioni poste a 3, 6 e 12 miglia nautiche dalla costa.

Le analisi dell'intero set di dati, che annovera oltre un migliaio di campioni, hanno mostrato preliminarmente un aumento del numero di *taxa* con oltre 350 record e hanno confermato la presenza di dinoflagellati potenzialmente tossici, come *Alexandrium minutum*, *Dinophysis caudata*, *Dinophysis sacculus*, *Lingulodinium polyedrum*, *Phalacroma rotundatum*, *Ostreopsis ovata*, *Prorocentrum lima*, *P. minimum* e varie specie di diatomee appartenenti al genere *Pseudo-nitzschia* già registrati prima del 2004. Inoltre è stata rilevata la presenza della rafidiofitea ittiossica *Chattonella subsalsa* e di altri dinoflagellati potenzialmente tossici, come *Dinophysis rapa*, *Gonyaulax spinifera*, *Karenia cf. brevis*, *K. cf. mikimotoi*, *K. cf. papillonacea* e *Protoceratium reticulatum*, osservati a partire dal 2005. Specie ritenute non indigene, come la diatomea *Pseudo-nitzschia multistriata* e il dinoflagellato bentonico *Ostreopsis ovata*, sono occasionalmente presenti nei campioni, quest'ultimo in corrispondenza di eventi di fioritura estiva. Infine, in relazione ai campioni prelevati nel monitoraggio per la strategia marina, sono stati rilevati, nelle stazioni più a largo, un numero ridotto di *taxa* e abbondanze minori con un contributo rilevante del gruppo atassonomico di nanoflagellati.

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Applicazione di oli essenziali come metodo non-invasivo per il controllo del biodeterioramento di beni culturali in pietra

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Il valore e la durabilità nel tempo dei beni culturali di tutto il mondo sono minacciati dal fenomeno del biodeterioramento, ovvero 'la perdita irreversibile di valore e/o informazione di un oggetto d'arte in seguito all'attacco di organismi viventi' come definito da Urzì e Krumbein (1994). In particolare, in presenza di luce naturale o artificiale e determinate condizioni ambientali, le superfici esposte di monumenti in pietra sono interessate dallo sviluppo di biofilm fototrofi. Questi biofilm sono comunità complesse immerse in una matrice polisaccaridica e formate principalmente da cianobatteri in associazione con batteri, microalghe, muschi e funghi, che sono in grado di formare patine colorate su vari tipi di substrato in ambienti aperti o confinati (Fig. 1) come cripte, chiese e ipogei (Albertano 2012, Rossi et al. 2012, Bruno, Valle, 2017).



Fig. 1
Monumenti in pietra interessati dal biodeterioramento: a) Complesso dei templi Lingaraj a Bhubaneswar, Orissa, India; b) Cubicolo illuminato da luce artificiale nelle Catacombe di Domitilla, Roma, Italia.

La presenza di questi biofilm sul bene culturale determina un danno estetico per l'alterazione dell'aspetto del bene stesso e a lungo termine può essere responsabile anche di danni strutturali che ne compromettono la stabilità e quindi la durabilità nel tempo. I metodi convenzionali utilizzati dai restauratori per l'eradicazione di questi biofilm dalle superfici di pregio consistono in spazzolature meccaniche accoppiate all'impiego di biocidi;

questi metodi risultano deleteri per i substrati, pericolosi per l'ambiente e dannosi per la salute umana (Urzì et al. 2016). Da qui la necessità di ricercare nuove soluzioni non-invasive per il substrato e anche eco-compatibili e non dannose per l'uomo (Bruno et al. 2014a, b, Hsieh et al. 2014, Bruno, Valle 2017, Ruffolo et al. 2017). Recentemente diversi oli essenziali, risultati non citotossici per le cellule umane e ampiamente impiegati per scopi medicinali (Gismondi et al. 2014, Giovannini et al. 2016), si sono dimostrati efficaci contro funghi e batteri presenti su beni culturali in pietra e legno (Stupar et al. 2014). Con l'obiettivo di sviluppare nuove strategie per il problema del biodeterioramento dei beni culturali e progettare un protocollo di facile utilizzo per il restauro dei patrimoni culturali in pietra, è stata quindi testata la potenzialità di due oli essenziali di poter essere utilizzati come 'biocidi verdi'. A tal fine sono state allestite piastre contenenti terreno minerale BG11 agarizzato (Rippka et al. 1979) su cui sono stati fatti crescere biofilm fototrofi costituiti da cianobatteri isolati dalle Catacombe Romane (Fig. 2).

Gli oli essenziali distillati da *Lavandula angustifolia* Mill. e *Thymus vulgaris* L. sono stati testati, singolarmente e in combinazione, in diverse concentrazioni, sui biofilm cresciuti su piastra.

L'effetto degli oli essenziali sui cianobatteri è stato monitorato mediante il controllo visivo della crescita del biofilm su agar e la valutazione della loro attività fotosintetica mediante il fluorimetro portatile mini-PAM. Poiché gli oli essenziali impiegati in questo studio erano di origine commerciale, i loro profili chimici sono stati studiati, in dettaglio, tramite gas cromatografia associata a spettrometria di massa (GC-MS). I profili GC-MS hanno mostrato come entrambi gli oli essenziali presentassero un elevato numero di metaboliti secondari vegetali dalle note proprietà antimicrobiche (come canfora, timolo) (Falcone et al. 2005, Chen et al. 2013). Al fine di valutare se questi oli essenziali potessero essere utilizzati anche su materiale di pregio senza indurre alterazioni al substrato, è stato anche valutato l'effetto della loro applicazione su pietre calcaree. I risultati ottenuti

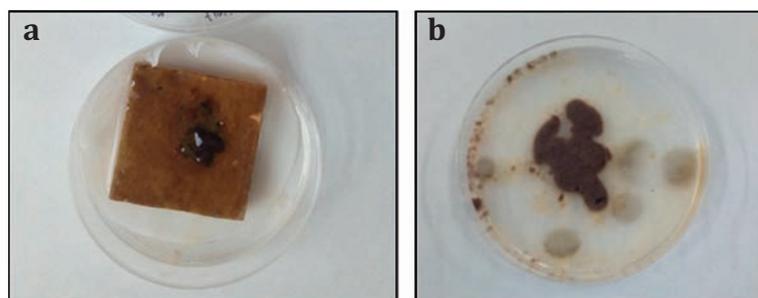


Fig. 2
I biofilm a cianobatteri utilizzati per testare l'efficacia degli oli essenziali come biocidi sono stati fatti crescere su a) pietre calcaree e b) piastre agarizzate.

Al fine di valutare se questi oli essenziali potessero essere utilizzati anche su materiale di pregio senza indurre alterazioni al substrato, è stato anche valutato l'effetto della loro applicazione su pietre calcaree. I risultati ottenuti

hanno evidenziato che gli oli essenziali erano in grado di ridurre efficacemente la crescita del biofilm in coltura diminuendo la loro attività fotosintetica; l'applicazione sulle pietre calcaree non ne alterava l'aspetto. In conclusione, lo studio ha dimostrato la fattibilità dell'impiego di oli essenziali come nuovi biocidi per il restauro di beni culturali.

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Harmful Algal dynamic in the gulf of Trieste during the last 30 years

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A 30-year (1986 – 2015) time series of data on phytoplankton abundances, collected in the Long Term Ecological Research (LTER) site in the Gulf of Trieste (North Adriatic Sea), has been analysed. A particular focus was put on the dynamics of potentially toxic species occurrence. The possible effects on economic activities based on marine living resources, such as mussel farming, a traditionally well-developed sector in the Gulf of Trieste, were also valued. During these decades several changes in phytoplankton abundances and communities composition were observed. An evident short-term as well as a interannual variability, with a range of more than one order of magnitude, characterized the 30 year – time of phytoplankton abundances in the Gulf of Trieste. The timing of interannual variability was different for total phytoplankton, diatoms and dinoflagellates, the latter two representing the main groups evaluated in this study. Total phytoplankton started to decline in the mid '90s, reaching the lowest abundances in 2006 and 2007, while a new increase was recorded after 2008, with several cases of abundances even higher than during the '80s and '90s. The diatoms time – series was characterised by a pronounced decrease in the mid '90s and by a prolonged one in the mid 2000, followed by an increase after 2009, with abundances exceeding 5 millions cells/l. Dinoflagellates abundances were characterised by a prolonged decrease from 2003 to 2010, followed by an increase in abundances up to values higher than during the beginning of the time series. The recent increase of dinoflagellates is of particular concern since the class includes also toxin-producing genera, which may adversely affect aquaculture and shellfish - farming activities located in the area. Considering dinoflagellates composition, some potentially toxic species were reported. Mostly of the recorded species belong to *Dinophysis* and *Alexandrium* genera, which caused contamination of mussels in the Gulf of Trieste since 1989. *Dinophysis caudata*, *D. sacculus*, *D. fortii*, undetermined *Alexandrium* species, *Gonyaulax polygramma* and *Lingulodinium polyedrum* are the most frequent taxa observed in all the samples collected. Considering the number of cases per year, both *Dinophysis* and *Alexandrium* genera increased in frequency after 2003. The highest frequency of *Dinophysis* occurrence resulted concomitant with episodic alerts due to the presence of biotoxins, leading to the bans of shellfish harvesting. Biotxin emergencies are among the more important threats to productivity indicated by local mussel farmers and the improved knowledge on their temporal trends and triggering factors are highly needed to protect public health and shellfish farming activities against economic losses.

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Dynamics of the phytoplankton communities in the Mar Piccolo of Taranto, an Italian Long-Term Ecosystem Research (LTER-Italy) site

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The Mar Piccolo of Taranto (Northern Ionian Sea), one of the most studied coastal areas in Italy, is a Long-Term Ecosystem Research site (LTER_EU_IT_095).

Phytoplankton data, both as active stages in the water column and cysts in the sediments, have been collected since 1991 (with gaps). Notwithstanding a reduction of nutrients caused by a relocation of wastewater inputs during the early 2000s, phytoplankton biomass (in terms of chlorophyll *a*) did not show significant variations over the years. However, other relevant changes occurred such as a reduction of the community size, due to a shift of dominance from diatoms to nanoflagellates, an increased duration of secondary blooms (beside the spring one), and the appearance of a diversified dinoflagellate community, comprised mixotrophic and heterotrophic species. An increase of the pico-sized planktonic component has also been recently detected. Moreover, wastewater diversion did not preserve the Mar Piccolo from harmful algal blooms and about 25 harmful species were monitored throughout the years.

These signals are in some ways mirrored in the sediments. In fact, the resting stages communities, during the last 20 years passed through many changes in their structure, even though calcareous dinocysts produced by Thoracosphaeraceae remained dominant. During the last 5 years, in particular, resting spores of some diatoms of the genus *Chaetoceros* were observed, together with the increase and new findings of some heterotrophic species, apparently a signal of the changes occurred in the Mar Piccolo. This change has been registered also with the disappearing of species, e.g. *Protoperidinium pentagonum* and *P. divaricatum*, whose cysts were regularly observed in surface sediments, even with low densities.

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Microphytobenthic community structure in four Po Delta lagoons subjected to natural and anthropogenic disturbance

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Benthic communities inhabiting the sediments of the Po Delta lagoons are subjected to multiple impacts. Besides those of natural origin, as the stress caused by the alternation of flood river conditions with the lean-season flow, several anthropogenic ones are present, e.g. excessive organic and nutrient loads deriving from urban and industrial discharges, and agricultural activities as well as high contamination levels. To investigate how the benthic communities respond to these disturbances, a scientific survey was carried out in May 2016 in the framework of the Project RITMARE. Four coastal lagoons of the Po River were chosen: two with more marine features (Scardovari and Caleri) and two more directly affected by the Po River freshwater (Canarin and Vallona-Marinetta). A greater freshwater input was mirrored in higher overall microphytobenthos (MPB) abundances in Marinetta-Vallona and Canarin (81600 ± 22735 and 81200 ± 50938 cells cm^{-3} , respectively) compared to Scardovari and Caleri (42350 ± 19131 and 48700 ± 10086 cells cm^{-3} , respectively). The MPB community was dominated by diatoms ($88.12 \pm 5.21\%$). In Marinetta-Vallona and Canarin, the higher relative abundance (RA) of Chlorophyceae, Cyanobacteria and freshwater planktonic diatoms was a clear signature of the major river inflow. Cluster and nMDS analyses highlighted different communities both at the intra-lagoonal level (among stations within the same lagoon), and inter-lagoonal level (among the four lagoons). Indeed, as revealed by diversity indices, the MPB community was more biodiverse at stations closest to the sea due to the concomitant presence of marine and brackish species, compared to the innermost stations where the lowest biodiversity was observed. Under organic enrichment conditions, as those encountered in Marinetta-Vallona due to intensive clam farming, *Nitzschia tryblionella* that prefers high organic loads was the most represented species (RA=19.1%). The benthic diatom community was also clearly affected by contamination levels. In Canarin, the most contaminated lagoon, diatom living forms that are scarcely or not at all associated with the sediments prevailed over the typical benthic forms, suggesting a negative influence attributed to contaminants accumulated in the sediments.

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Flora marina bentonica del Mediterraneo: Rhodophyta (*Rhodymeniophycidae* escluse)

M. Cormaci, G. Furnari, G. Alongi

Dopo una breve introduzione sui caratteri generali delle Rhodophyta (Alghe Rosse) e sulla loro classificazione, distribuzione geografica ed ecologia, viene riportata una chiave d'identificazione dei generi e dei *taxa* specifici e infraspecifici macroalgali di questo Phylum [ad esclusione delle *Rhodymeniophycidae* che saranno oggetto di altro contributo] presenti in Mediterraneo [Mar Nero escluso].

A seguito di una revisione tassonomico-nomenclaturale, 120 *taxa* a livello specifico e infraspecifico sono stati accettati e trattati. 44 generi, 15 famiglie, 12 ordini, 3 sottoclassi, 4 classi e 2 subphyla sono rappresentati. Inoltre 61 *taxa*, a livello specifico e infraspecifico, sono stati considerati *taxa excludenda*, di cui 18 da confermare in Mediterraneo; 90 *taxa*, a livello specifico e infraspecifico, e 1 a livello di genere, sono considerati *taxa inquirenda*; 29 *taxa*, a livello specifico e infraspecifico, sono risultati nomi illegittimi; 33 *taxa*, a livello specifico e infraspecifico, e 1 a livello di famiglia, sono risultati invalidi; infine sono state individuate 17 combinazioni invalide, mentre vengono proposte 4 nuove combinazioni e un nome nuovo.

L'aggiornamento tassonomico segue i più recenti lavori basati sull'analisi molecolare. Di ogni *taxon* viene data una breve descrizione preceduta da alcuni riferimenti bibliografici riportanti sue illustrazioni e/o la sua distribuzione in Mediterraneo. Inoltre, la trattazione della maggior parte dei *taxa* è arricchita da note a supporto delle sinonimie indicate, o delle scelte tassonomiche seguite, o delle motivazioni per cui il *taxon* è stato considerato *inquirendum* o *excludendum*. Il lavoro è completato da un glossario di 148 voci, da un indice di tutti i *taxa* citati nel testo e da 52 tavole.

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Tracking the arrival and spread of non-indigenous macroalgae in New Zealand

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New Zealand has an effective Biosecurity System that helps to protect New Zealand's economy, environment and human health from harmful organisms like pests and diseases. In the marine environment, control of pests/diseases is problematic and preventing the introduction of non-indigenous species is very difficult. Nevertheless, since 2002 New Zealand's Ministry for Primary Industries (MPI) has funded a national program of marine surveillance at the 11 most heavily-used ports and marinas to detect non-indigenous species. Approximately 3000 survey locations at these 'high-risk' sites are checked twice a year, through a variety of survey methods. Suspected non-indigenous specimens, or organisms whose identity is unknown, collected during these surveys are sent to the Marine Invasives Taxonomy Service (MITS) for taxonomic identification. The Biosecurity System depends on the ability to distinguish between indigenous and non-indigenous species. The identification of macroalgae is particularly challenging as the New Zealand flora is still far from being fully documented. Species of macroalgae were being recognized as introduced to New Zealand waters from the 1970s onwards, and in 1983, Adams assembled the first checklist of 14 marine macroalgae that she hypothesized were non-native species, based on their distribution patterns. Currently, 48 algal species are considered non-native to New Zealand (13 brown, 23 reds and 12 green algae), with about half of these being recognized as such in the last decade. Some of these introduced species have a restricted distribution in New Zealand while others (e.g., the Asian kelp *Undaria pinnatifida* (Harvey) Suringar or *Grateloupia turuturu* Yamada), are widespread. Particularly challenging are species belong to *Polysiphonia sensu lato* or *Ulva* spp. that are common on vessels, and require DNA analysis for precise identification. Some examples of past and current research on macroalgae will be presented.

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A journey of discoveries: New Zealand Kallymeniaceae

R. D'Archino

The seaweeds of New Zealand have always challenged taxonomists. Part of the problem is that concepts and names of algae from other parts of the world have been used. Also New Zealand includes diverse geographic regions from the subantarctic islands to subtropical waters. Over the last 10 years we have realized that the Kallymeniaceae of New Zealand are more diverse and unique than we ever could have imagined. New genera have been established, *Psaromenia*, *Stauromenia*, *Judithia*, *Wendya*, *Blastophyllis* and *Zuccarelloa*, to accommodate species previously assigned to other genera, but the great challenges are still ahead; the delimitation of the genus *Callophyllis* and the increasing number of new discoveries. The number of taxa increased from 14 in 2007 (6 genera) to 18 in 2017 (9 genera), but, including the undescribed taxa, currently we have identified 37 species (23 genera). Some species names have been used largely in the past eg. *Callophyllis ornata* rarely collected from its type locality - Auckland Is, but this name has been applied to numerous specimens with wide ranging morphologies. Kallymeniaceae are notoriously difficult to identify as critical distinguishing characters frequently are based on fertile material at the right stage of maturity. Often our samples are sterile or come from remote locations (eg. subantarctic islands, Three Kings, deep water), where only occasional collections have been made, and for some of these undescribed taxa only few specimens are available. Recently, great diversity has been uncovered in Australia, increasing the number of Kallymeniaceae genera to 43, and while it was expected these two countries would share common taxa, it seems that New Zealand has very unique Kallymeniaceae.

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A multi-gene phylogeny to uncover species diversity in the planktonic diatom genus *Chaetoceros* Ehrenberg, 1844

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The genus *Chaetoceros* (Ehrenberg, 1844) is arguably the most species-rich among planktonic diatoms, with more than 200 taxonomically accepted species distributed in upwelling regions and coastal areas all over the globe. Its species are recognized easily because they possess hollow spines, called *setae*. However, species identification is less easy since morphology is variable and affected by culture conditions and environmental factors. Previous analyses resulted in the characterization of hundreds of *Chaetoceros* strains by means of morphological (light, scanning and transmission electron microscopy) and genetic (D1-D4 region of 28S rDNA, 18S rDNA) data. Results show that several terminal clades (operational taxonomic units or species) are still unknown to science and that many morphologically defined species actually consist of species complexes (e.g. *C. affinis*, *C. curvisetus*, *C. lorenzianus*, *C. socialis*). Here we present a multi-gene phylogeny of *Chaetoceros* using nuclear (18S and 28S rDNA), chloroplast (*rbcL* and *psbA*) and mitochondrial (COI) genes to assess if the signal obtained from the evolutionary history of organelle genes is congruent with the nuclear ones. Preliminary results show that the phylogenies are, indeed, congruent. Some discrepancies exist in species complexes, with *rbcL* being more effective in discriminating closely related species than *psbA*. Also bootstrap support to internal nodes was higher for *rbcL* than for *psbA*, probably because of a higher substitution rate of the former gene. The COI tree topology differed markedly from that of the other gene trees, mostly due to the high substitution rate of the marker. The latter aspect makes this gene region a good marker for population level inferences or barcoding approaches.

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Survey on dinoflagellate cysts in recent marine sediments of the port of Trieste (Northern Adriatic Sea)

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In coastal and temperate shallow waters, dinoflagellates can produce benthic resting stages to overcome unfavorable periods, and germinate in the water column once favorable environmental conditions are re-established. Studies on the composition, abundance and distribution of cysts in surface sediments of semi-enclosed bays such as ports can provide valuable information both to detect potential harmful and non-indigenous species and to foresee dinoflagellate dynamics inside the phytoplankton community. To this aim, a monitoring programme was carried out in the port of Trieste within the framework of the European IPA project BALMAS (Ballast Water Management System for Adriatic Sea Protection). In May 2014 and March 2015, 4 sites (Ts1-2-3-4) characterized by maritime traffic and anthropic pressures and a reference site (C1) located in the Marine Reserve of Miramare (Gulf of Trieste) were sampled. Surface sediments were collected by means of a box corer and samples consequently processed. The identification of the resting stages was based on the shape and morphology of cyst bodies (Matsuoka, Fukuyo 2000) using the inverted microscope at 20x objective. Furthermore, to verify the taxonomic identification, some selected cysts were isolated and induced to germinate.

In this study, a total of 38 dinoflagellate taxa were recognized. Cyst assemblage resulted dominated by *Gonyaulacales* and calcareous *Peridinales*, mainly represented by the potentially toxic *Lingulodinium polyedrum* and by resting cysts of the genus *Alexandrium* in the size 20 μm . In May 2014 cysts varied from a minimum of 17 to a maximum of 232 cells g^{-1} of dry weight detected respectively at C1 and Ts4. In March 2015, cysts ranged from 21 to 60 cells g^{-1} of dry weight respectively at Ts2 and Ts4. The latter site Ts4 recorded the higher cyst deposits detected in the port. From the germination experiments carried out on the cysts collected during the second survey (March 2015), a new record for the northern Adriatic, the species *Alexandrium margalefii*, was developed. Other potentially toxic morphotypes of the genus *Alexandrium* identified in both campaigns were already renowned in this area. Compared to previous few studies conducted in the Gulf of Trieste, the number of taxa and the most representative morphotypes were comparable to past studies whereas abundances were much lower than past monitoring. Our results remark the importance of these stages of dinoflagellate life cycle, and the necessity to intensify investigation especially in areas at high risk such as ports.

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The potential role of phytoplankton in traps of carnivorous bladderworts (*Utricularia*, Lentibulariaceae)

N.T.W. Ellwood, R. Congestri, S. Ceschin

Carnivorous aquatic plants of the genus *Utricularia* (Lentibulariaceae) are well known to capture small aquatic invertebrates, using structures called bladders. But information is coming to the fore suggesting that *Utricularia* could be re-classified as an omnivore as phytoplankton species can make up a major proportion of bladder contents. It is not fully understood if phytoplankton cells are preyed upon or if they are taken in merely as a consequence when invertebrate prey triggers the bladder. Past studies have indicated that both scenarios could be possible. Some studies now also suggest that a mutual-symbiotic relationship exists between *Utricularia* and the community in the within the bladders. The present study is to determine a potential role for phytoplankton in nutrient supply to *Utricularia*.

The identification and quantification of phytoplankton species were carried out on the contents of bladders sampled from the carnivorous bladderwort *Utricularia australis*. Samples of three full *Utricularia* filaments were taken from 18 central Italian sites and the contents of 10 or 15 randomly chosen bladders were observed. These data were compared to the external phytoplankton community and water chemico-physical and nutrient variables. Every bladder was shown to contain some phytoplankton cells, with findings showing species spread across 6 divisions (Chlorophyta, Bacillariophyta, Pyrrophyta, Euglenophyta, Cryptophyta, Cyanophyta). Significant relationships of phytoplankton species assemblages within the bladders and those in the ambient water were found. Also in the majority of bladders there were a high proportion of live compared to dead cells. Many bladders had phytoplankton-only communities. It was not possible to say for certain if phytoplankton was preyed upon or not, or if there was a mutualistic-symbiotic relationship, but the presence of many living cells could suggest the latter option is possible. The findings here imply that the bladder phytoplankton community is a consequence of the ambient community given the relationships between them. The presence of phytoplankton-only contents in many bladders implies possible automatic - rather than prey - triggered bladder functioning.

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Gene expression changes in diatoms in response to external stimuli

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Signal transduction mechanisms are still poorly defined in marine phytoplankton, including diatoms. Yet, understanding how cells sense and integrate environmental signals is fundamental to explain population dynamics, community interactions and ecosystem functioning. We use several approaches, mostly based on transcriptomics and loss of function strategies, to study molecular changes occurring in diatom cells upon reception of external stimuli, such as changes in turbulence, or chemical cues produced during sexual reproduction or grazing, using dedicated, custom-made instrumentation.

I will present data on diatom responses to turbulence. It is known that diatoms thrive in turbulent environments, however, despite several experimental and numerical studies, if and how diatoms may profit from turbulence is still an open question. One of the leading arguments is that turbulence favours nutrient uptake. To study this process, we designed and built the TURBOGEN, a prototypic instrument that can generate natural levels of microscale turbulence. We tested several diatom species using TURBOGEN, and demonstrated that diatoms actively respond to turbulence in non-limiting nutrient conditions by tuning their chain length. Differential expression analyses were conducted to study the transcriptional response to turbulence of *Chaetoceros decipiens*, one of the species displaying obvious morphological changes. *C. decipiens* responded to turbulence by activating energy storage pathways like fatty acid biosynthesis. In addition, in experiments lasting 12 days, we observed that in turbulence *C. decipiens* continued to take up phosphorus and carbon even when silicon was depleted. These findings indicate that turbulence affects diatoms in a more sophisticated fashion than what was accepted so far.

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A putative phytochelatin is involved in Cr(VI) tolerance in strict relation with sulphur metabolism in *Scenedesmus acutus* (Chlorophyceae)

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Metal contamination is a serious environmental problem that affects life forms and changes the natural microbiota of aquatic ecosystems.

Chromium (Cr) is one of several heavy metals (HM) that causes serious environmental contamination in soil, sediments, and groundwater. Cr can exist in the environment as Cr(III) or Cr(VI), and particularly Cr(VI) form is extremely toxic, mutagenic, and carcinogenic.

Microalgae resistance and/or detoxification to heavy metals may occur through several mechanisms: binding to the cell wall; low plasma membrane permeability; active extrusion; biotransformation; compartmentalization of HM into vacuoles and other intracellular organelles; complexation with chelating agents, such as non-proteinaceous compounds (as malate, citrate, and polyphosphates) or metal-binding proteins, such as metallothioneins and phytochelatins. Phytochelatins (PCs) are heavy metals chelating cysteine rich polypeptides with the general structure $(\gamma\text{-Glu-Cys})_n\text{-Gly}$. Phytochelatins are produced by the enzyme Phytochelatin Synthase (PC-synthase) which uses glutathione (GSH) as its substrate. It is now well established that glutathione (GSH) and PCs are a central component of the plant defence system against various forms of stresses.

Several physiological studies in plants, indicated the role of PCs in the homeostasis and detoxification of toxic metals including Cr. However, synthesis of PCs has received little attention in algal cells and no data regarding Cr-induced PCs synthesis in microalgae exist.

In this work, we reported the first evidence of a PCs synthase gene from the microalga *Scenedesmus acutus*. Using degenerate primers, we amplified a partial cDNA fragment of nearly 1300 bp (*SaPCS*). The deduced protein shows a high identity homology (46%-74%) with other PCs synthase of algae and plants.

In order to define its role in Cr detoxification, we analysed by RT-PCR the level of *SaPCS* transcription in two strains of *S. acutus* with different chromium sensitivity (Cr tolerant strain vs wild type) after 24h culture in standard and in Cr supplemented medium (1mg/l Cr (VI)). Since the pathway of Cr (VI) translocation is an active mechanism involving transporter of essential nutrients such as phosphate and sulphate, we also evaluated the role of PCs synthase in the two different *S. acutus* strains after sulphur starvation.

Our preliminary data indicate that in the Cr-tolerant strain *SaPCS* transcription is induced following Cr(VI) treatment both in S-sufficient and in S-replete condition whereas in the wild type the induction is significantly enhanced by sulphur starvation. These observations indicate that in *S. acutus* PC synthase is not only involved in Cr detoxification, but can also play a role in the balance the intracellular sulphur.

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Monitoring the cyanobacteria species biodiversity and growth during the maturation of Euganean Thermal Muds

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The Euganean Thermal District is the oldest and largest thermal center in Europe and represents a reference in mud-based thermal treatments. The mature mud is the result of traditional methods, handed down over the centuries up to our days. During the mud maturation process, we can observe the formation of a biofilm, mainly consisting of cyanobacteria. They are suggested to produce active molecules conferring the therapeutic properties of the mud. Among them, *Phormidium* ETS-05, an endemic species firstly isolated from the Euganean District, is well known for the production of galactosyldiacylglycerols with anti-inflammatory effects (Bruno et al. 2005, doi: 10.1016/j.ejphar.2005.09.023), leading to the achievement of the European Patent on the active principles of the Euganean Thermal Muds in 2013 (EP1571203). Recently, another endemic species, *Cyanobacterium aponinum* ETS-03, has been studied for its ability to produce polysaccharides with immunostimulating effects (Gudmundsdottir et al. 2015, doi: 10.1016/j.imlet.2014.11.008; Gris et al. 2017, doi: 10.1007/s10811-017-1133-3). A raw protocol, called "Disciplinare per la Tutela del Marchio Collettivo di Origine del Fango del Bacino Termale Euganeo" has been written to formalize the traditional methods. Therefore, the scientific bases and a deep knowledge of all elements influencing the mature mud formation are still missing.

In the first part of this study, the maturation process of 2 thermal SPAs has been monitored every 15 days for 2 months, following the cyanobacteria population growth and composition as function of time and environmental parameters. In the second part, the presence and the total amount of the Patent target species, *Phormidium* ETS-05, in mature muds of 33 different thermal SPA, has been investigated as an indicator of the correct maturation process.

Results showed that, during maturation, the maintenance of temperature in a proper constant range is one of the crucial aspects to consider to have a proper final mature mud. The observations and the analysis of mature muds of different SPAs revealed that the temperature influences both the final concentration and the cyanobacteria population composition. The target species *Phormidium* ETS-05 was present in almost all the monitored SPAs, and represented always at least the 40% of the total cyanobacteria population.

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Ten years of research on *Ostreopsis cf. ovata* (Dinophyceae) in the Gulf of Trieste: an interdisciplinary approach

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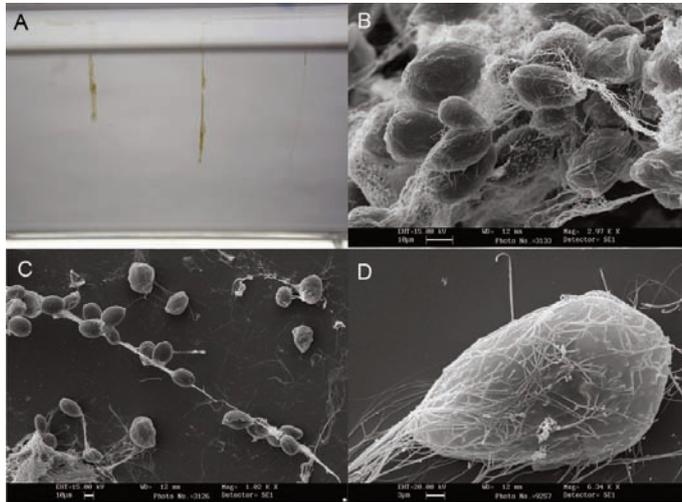


Fig. 1
Ostreopsis cf. ovata cultures from the Gulf of Trieste. A. Cells in a culture flask forming filamentous floating aggregates departing from water surface. B. Scanning electron microscope (SEM) image of cell aggregates with high amounts of mucilage formed by filaments and amorphous material. C. SEM image of a mucilage filament connecting more cells together. D. SEM image of a cell extruding numerous trichocysts joining together at the ventral end of cell.

omy to cell ultrastructure, genetic and molecular characterization, toxin profile, intracellular localization of toxins and other metabolites by immunochemistry and/or Raman spectroscopy. A short review of the results obtained is given.

Taxonomic identification: *Ostreopsis* cells of the Gulf of Trieste were identified as *Ostreopsis cf. ovata* by light microscopy and molecular analysis. Final alignment of *Ostreopsis* sequences with the Genbank sequence database of *Ostreopsis* species confirmed that the *Ostreopsis* isolates belonged to the *O. cf. ovata* Atlantic/Mediterranean clade (Honsell et al. 2011).

Cell structure: new ultrastructural features were described by scanning and transmission electron microscopy. *Ostreopsis* mucilage appeared to be formed by a network of trichocysts extruded through thecal pores and by amorphous polysaccharidic material deriving from mucocysts (Fig. 1). A role of trichocysts as fibrillar component of mucilage seems to be a specific feature of *Ostreopsis*, not observed in other benthic dinoflagellates like *Coolia monotis* or *Prorocentrum lima*. Two different types of plastids were found in all stages of cell growth: large elongated chloroplasts with stacked thylakoids in peripheral cytoplasm and small round proplastid-like structures in the central part of the cell. A considerable part of cytoplasm was occupied by neutral lipid droplets. Raman spectroscopy showed also the presence of poly-unsaturated lipids, concentrated in spots at the cell border (Honsell et al. 2013).

Toxin analysis: several palytoxin-like compounds (ovatoxin-a,-b,-c,-d,-e) were identified by LC- HRMS in natural populations, ovatoxin-a being the most abundant (Honsell et al. 2011). In culture, the highest toxin content was reached during the senescent phase (Honsell et al. 2013).

Immunocytochemistry: the presence of PLTX-like compounds in *O. cf. ovata* cells was investigated by immunocytochemistry, using monoclonal or polyclonal anti-PLTX antibodies. Both antibodies reactions localized PLTX-like compounds in a cytoplasmic network made of filaments and small dots surrounding chloroplasts and other organelles, and in numerous brighter dots in the peripheral cytoplasm (Honsell et al. 2011).

Detection methods: a panel of different detection methods was set up and evaluated for their suitability to quantify PLTXs in microalgal samples and mussels. A LC-HRMS method was developed for identification, structural characterization and quantitation of the entire array of toxins produced by *O. cf. ovata* in algal pellets (Ciminiello et al. 2012), seawater, and marine aerosols (Tartaglione et al. 2016). Combination of solid phase extraction with

Since 2000s, benthic blooms of the harmful dinoflagellate *Ostreopsis cf. ovata* have been occurring with increasing frequency and extension along the Mediterranean coasts. Concomitantly to these events, episodes of human adverse effects associated mainly with cutaneous and inhalational exposure to marine aerosols and/or seawater have been recorded. Although high levels of *Ostreopsis* toxins have been detected in edible marine organisms, no human cases of seafood poisoning ascribed to these compounds have been documented, so far. Mediterranean strains of *O. cf. ovata* have been found to produce palytoxin (PLTX) analogues, with ovatoxin-a (OVTX-a) as the major toxin, in addition to other ovatoxins and low amounts of isobaric palytoxin. In the Gulf of Trieste, *O. cf. ovata* was first recorded in 2006, but the real beginning of an extensive research on this organism dates back to 2009 when a bloom occurred near the small beach of Canovella de' Zoppoli (Trieste). This improved the collaboration within a research team of biologists, chemists and toxicologists for an interdisciplinary study providing new insights on different features of this organism: from taxon-

LC-HRMS detection proved able also to determine the presence of ovatoxins in mussels at level as low as 15 g/kg, well below the challenging maximum permitted level of PLTX recommended by EFSA of 30 g/kg (Ciminiello et al. 2015). An indirect sandwich ELISA was initially developed using murine monoclonal and rabbit polyclonal antibodies against PLTX. The good sensitivity (limit of detection, LOD, and quantitation, LOQ, of 1.1 and 2.2 ng/mL, respectively), its good accuracy and reproducibility demonstrated its suitability to detect PLTX, also in contaminated mussels (LOQ = 11 µg/kg) below the limits suggested by the European Food Safety Authority (Boscolo et al. 2013).

The sensitivity of the sandwich ELISA was further improved by an ultrasensitive electrochemiluminescence-based sensor, taking advantage of the specificity provided by anti-PLTX antibodies, the good conductive properties of carbon nanotubes, and the excellent sensitivity achieved by a luminescence-based transducer. The sensor was able to produce a concentration-dependent light signal, allowing PLTX quantification in mussels, with a LOQ of 2.2 µg/kg of mussel meat, more than 2 orders of magnitude more sensitive than that of the commonly used detection techniques, such as LC-MS/MS (Zamolo et al. 2012).

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Physiological adaptation within the Chaetoceraceae to the ecological challenges in the northern Adriatic

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The northern Adriatic is a suitable ecosystem for observing phytoplankton under a variety of different ecological conditions. This shallow basin is characterized by a multitude of steep and quickly changing gradients (e.g. nutrients), both temporal and spatial. The current systems and the major freshwater input (River Po) generate strong gradients in nutrient concentrations with an expressed N/P imbalance and sustained phosphate limitation. The genus *Chaetoceros* is among the most diverse and species rich diatoms often dominating the phytoplankton community in the Adriatic Sea. There is a big number of species belonging to this genus with different physiology, seasonality and ecology.

We analyzed species from the genus *Chaetoceros*, with a special focus on their metabolic reaction towards phosphate limitation. Here we report data on species-specific growth rates under different nutrient regimes, phosphate uptake rates, alkaline phosphatase activity, localization and activation patterns and characteristics of alkaline phosphatase activity. Our results demonstrate a high interspecific variation in metabolic responses to phosphate limitation in sympatric congeneric species. Ecological characteristics and hence significance and function appears hence to be defined on the species level and appears far from homogeneous within genera. This results in a highly structured planktonic ecosystem that allows for a high level of sympatric congeneric species diversity.

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Citizens and scientists work together to monitor marine alien macrophytes

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The introduction of non-indigenous species (NIS) is an ongoing phenomenon which has been pointed out as a major threat to biodiversity at different levels (Wallentinus, Nyberg 2007, Katsanevakis et al. 2014, Vergés et al. 2016). NIS may in time become invasive (Invasive Alien Species “IAS”) and may cause biodiversity loss and ecosystem service changes (Brunel et al. 2013, Giakoumi 2014, Vergés et al. 2016). The Mediterranean Sea is an important hotspot for marine NIS (ca. 1,000 such species recorded to date, Zenetos et al. 2012, Galil et al. 2015, Verlaque et al. 2015). To reduce the risk of future IAS introduction and to better understand their invasive potential and spread dynamics, monitoring and surveillance plans are required. The creation of permanent alarm systems and public awareness campaigns are crucial for reducing the risk of IAS introduction. Since intensive monitoring programs could be very expensive, citizen science, involving citizens (e.g. tourists, fishermen, divers) in the collection of data, could be a useful tool for providing data on IAS, that would otherwise be impossible to collect because of limitations on time and resources. Citizen science is having an increasing success worldwide. Citizen science projects has rapidly and enormously increased in recent years (Conrad, Hilchey 2011), also thanks to the wide availability of mobile technologies and internet access that enable an easy and cheap way to communicate, share and interchange data. The value of citizen science has been widely recognized. Of course, in order to be used for scientific purposes and management decisions, the collected data need appropriate quality assurance measures such as validation and verification by taxonomic experts. We report on the experience of two citizen science projects: the Project “*Caulerpa cylindracea* – Egadi Islands” and the Project “Invasive Algae”, included within the “Seawatchers” platform. The first one, sponsored by the STEBICEF Department of the University of Palermo and by the Egadi Islands Marine Protected Area (MPA), aimed at creating a database on the spread dynamics of *Caulerpa cylindracea* Sonder within the Egadi Islands MPA (western coast of Sicily, Tyrrhenian Sea). Among IAS, *C. cylindracea*, introduced from Australia and New Caledonia (Belton et al. 2014), has raised serious concern due to its ascertained impact on Mediterranean communities (Klein, Verlaque 2008, Papini et al. 2013, Katsanevakis et al. 2014). The project, presented during the International Congress GeoSub2016 (Mannino et al. 2016), allowed to gather 156 sightings, mainly recorded by citizens (Fig. 1). Useful information on the behaviour strategies of the alga was also collected. The second one, coordinated by the Institute of Marine Sciences of Barcelona (CSIC, Spain), collects data on 10 marine IAS (*Acrothamnion preissii* (Sonder) E.M. Wollaston, *Asparagopsis armata* Harvey, *A. taxiformis* (Delile) Trevisan, *C. cylindracea*, *C. taxifolia* (M. Vahl) C. Agardh, *Codium fragile* subsp. *fragile* (Suringar) Hariot, *Halimeda incrassate* (J. Ellis) J.V. Lamouroux, *Lophocladia lallemandii* (Montagne) F. Schmitz, *Styopodium schimperi* (Kützinger) Verlaque & Boudouresque, *Womersleyella setacea* (Hollenberg) R.E.Norris). These projects highlight how important the contribution of citizen science campaigns is for collecting new data and information on marine NIS and to significantly improve the efficacy of monitoring and surveillance plans. Moreover, in areas particularly vulnerable to biological invasions, such as Sicily, they represent an opportunity to promote the creation of early-warning systems, and an effective tool in the management of present and future introductions of NIS within the Mediterranean Sea.

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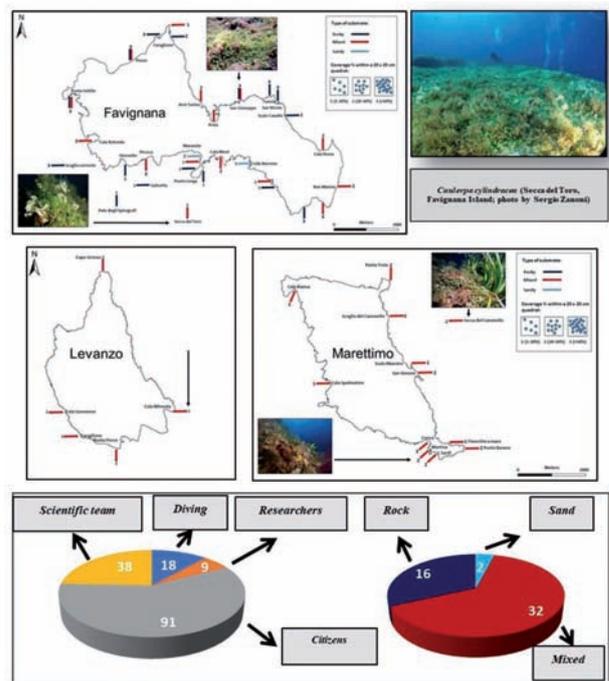


Fig. 1
Main results of the project “*Caulerpa cylindracea* – Egadi Islands”: sightings recorded in the main islands, groups of volunteers involved in the project, substrates colonized by *Caulerpa cylindracea*.

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***In vitro* isolation of anti-*Alexandrium minutum* nanobodies from a pre-immune library**

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At the present, the identification of planktonic species in coastal water mostly relies on light microscopy observations, only in some cases through PCR analysis and for specific morphological analyses by electronic microscopy. The analyses are performed by highly trained personnel and require lab equipment and long time in order to get the results. The increasing danger represented by Harmful Algal Blooms (HABs) and the new legislation on ballast water management urge faster and possibly easy-to-perform diagnostic methods able to guarantee sensitivity and reliability. Immuno-reagents for microalgae capture would have the requisites of specificity necessary for developing semi-automatic sensitive biosensors to be operated on-site by generic personnel. This alternative is at the moment neglected because of the elevated costs for monoclonal antibody selection and production.

Here we demonstrate for the first time the feasibility to recover nanobodies (VHHs) selective for native surface epitopes of *Alexandrium minutum* by direct whole cell biopanning using a pre-immune phage display library. The recombinant nature of VHHs enabled their bacterial expression as eGFP fluorescent reagents. We demonstrated that these reagents are directly suitable for fluorescence microscopy and flow cytometry, but could be used also on solid surfaces such as ELISA plates. The isolated fluorescent nanobodies showed no cross-reactivity with other microalgae of the Collection of Sea Microorganisms (COSMI) available at INOGS such as *Scrippsiella lachrymose* (COSMI n: 1049), *Prorocentrum minimum* (COSMI n. 1034), *Pleurochrysis cf. roscoffensis* (COSMI n. 5005), Cryptophycean *sp. 2* (COSMI n. 3002) and *Phaeodactylum tricornutum* (COSMI n. 2007). They were able to bind selectively the target cells in both fixed and fresh samples with minimal processing, demonstrating the advantage of these reagents over conventional IgGs. The establishment of the protocol represents a proof-of-principle that can be applied to further species. It is possible to isolate monoclonal nanobodies in 1:3 of the time and at a cost roughly 1:10 of that necessary to select a monoclonal antibody by hybridoma technique. The production costs are between 10 and 100 times lower. Furthermore, nanobodies are easy to engineer and, therefore, optimal material for biotechnological applications (functionalization of solid surfaces, nanoparticles, sensor chips).

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Assessment of *Posidonia oceanica* (L.) Delile (1813) meadows along the Italian coasts: an excursus of 25 years

C. Micheli

The aim of this study was to estimate time series of data of *Posidonia oceanica* (L.) Delile (1813) meadows growing along the Italian islands and in the northern-central Tyrrhenian coasts, during 25 years (1991-2016) of monitoring studies.

During the years we have assessed the ecological status of the species integrating the knowledge acquired about genetic, biomass, density and morphological features of the populations in relation to different natural and anthropogenic impacts and various substrate composition (rocky, sand and mat) (Micheli et al. 2018). *P. oceanica* meadows growing along the coastline are under continuous natural and anthropogenic pressure where the way meadow health correlates with its genetic and genotypic diversity (Micheli et al. 2005, 2015).

Previous studies have revealed the important role of currents (Micheli et al. 2010a, Rotini et al. 2011) and during 2003–2004, when a massive fruiting event occurred; fruits were collected from plants at Monterosso al Mare (meadow) and compared with stranded fruits sampled in front of the meadow and down current in Tuscany along 80 km of the coast (Aliani et al. 2006, Micheli et al. 2010a). After their growth in culture, the plants were analyzed by RAPD molecular markers. Cluster analysis of similarity showed four distinct genetic populations revealing their provenience.

Moreover, due to different anthropogenic impacts, such as a nourishment work, changes in the ecological structure of the *P. oceanica* meadows, at Monterosso a mare (Ligurian Gulf) were observed over ten years (Micheli et al. 2012).

Recently we have identified new diagnostic tools for effective monitoring of the conservation status of *P. oceanica* beds by applying innovative methodologies like remote sensing techniques (Micheli et al. 2010b, Borfecchia et al. 2013a,b), and using a multidisciplinary method as a suite of indicators to highlight the seasonal variation of biomass, phenols and the main biotic parameters as well as pigment concentration of seagrass meadows (Rotini et al. 2013).

Knowledge of the ecological structure of meadows is considered a pre-requisite for many multidisciplinary programs as well as the marine energy projects by which the renewable wave energy resource can be exploited in the Mediterranean (Borfecchia et al., 2016 Micheli et al. 2017).

Assessment of *P. oceanica* meadows along the coasts allowed us to evaluate the role of the species in sustaining the primary production and the trophic level in the Mediterranean.

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Towards the assessment of the diversity of the dinoflagellate genus *Tripos*: single cell approach

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The dinoflagellate genus *Tripos* (previously included in *Ceratium*) represents a significant component of marine microplankton. The genus includes more than 120 morphologically described species and infraspecific taxa, such as forms and varieties, found from polar to tropical areas. The genus has been suggested as indicator of ocean warming because of its taxonomic richness, wide geographical distribution and geographical patterning of its taxa related to temperature. However, the great variability of shapes described even within a single strain, coupled with the lack of clear discriminating characters, makes challenging species circumscription based on morphological criteria.

Using a single-cell approach, 52 *Tripos* cells isolated from the Gulf of Naples were characterized from the molecular (28S and 18S) and morphological point of view. In most cases, sequences obtained from morphologically identical cells clustered in the same clade. The 28S phylogeny showed a better capacity to discriminate different species than 18S, but also suggested the existence of cryptic diversity in the genus. As an example, cells morphologically identified as *T. trichoceros* clustered into two different clades.

Taxonomically validated sequences were added to the existing reference database DinoREF to improve the interpretation of metabarcoding data from a time series collected in the Gulf of Naples (LTER MC, 48 dates in 3 years).

I will present preliminary results and discuss problems and future possible approaches to study the diversity of this important but poorly known genus.

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BiOLEAP Biotechnological optimization of light use efficiency in algae photobioreactors

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New renewable energy source are highly needed to compensate exhausting fossil fuels reserves and reduce greenhouse gases emissions. Some species of algae have an interesting potential as feedstock for the production of biodiesel thanks to their ability to accumulate large amount of lipids. Strong research efforts are however needed to fulfil this potential and address many issues involving optimization of cultivation systems, biomass harvesting and algae genetic improvement. This proposal aims to address one of these issues, the optimization of algae light use efficiency. Light, in fact, provides the energy supporting algae growth and must be exploited with the highest possible efficiency to achieve sufficient productivity and make their cultivation competitive. Algae efficiency in converting solar radiation, however, depends on many environmental factors, including light intensity, temperature, nutrient and CO₂ availability. Optimizing microalgae productivity in such a complex environment hinges on our ability to describe, in a quantitative manner, the effect of these various parameters as well as their mutual interactions. Application of computational models that are capable of quantitative predictions can prove especially useful in identifying which parameters have the largest impact on productivity, thereby providing a mean for enhancing growth through design and operational changes. This information has now been employed to select *Nannochloropsis gaditana* strains with altered regulation of photosynthesis that showed improved productivity in lab scale photobioreactors.

Light influence on algae metabolism has also been investigated using a combination of genomic, transcriptomic and metabolomics analyses. These evidenced how light availability in *N. gaditana* modulates the carbon partitioning and TAG biosynthesis by affecting the transport of reduced carbon in and out of the chloroplast.

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Phytoplankton diversity in Adriatic ports

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Besides being a highly sensitive ecosystem, the Adriatic Sea is also an important seaway for international and local shipping, which poses a serious risk on the transfer of harmful aquatic organisms and pathogens (HAOP) by ships. Monitoring programs and surveillance activities would be of limited efficacy if carried out on a local scale only. The first step in setting up a common Adriatic approach towards unwanted threats to the environment from the transfer of HAOP was to prepare an inventory of organisms in ports. Phytoplankton diversity was determined through the port baseline survey in twelve Adriatic ports in years 2011, 2014 and 2015. Phytoplanktonic HAOP were identified as species, either native or non-indigenous (NIS) that can trigger harmful algal blooms (HAB). A list of 689 taxa was prepared and among them 52 were classified as HAB and five as NIS. Records of toxigenic NIS (*Pseudo-nitzschia multistriata*, *Ostreopsis* species including *O. cf. ovata*) indicate that the intrusion of non-native invasive phytoplankton species has already occurred in some Adriatic ports. The seasonal occurrence and abundance of HAOP offers a solid baseline for a monitoring design in ports in order to prevent possible expansion of HAOP outside their native region.

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Different features of the ultrastructure of autophagy in algae

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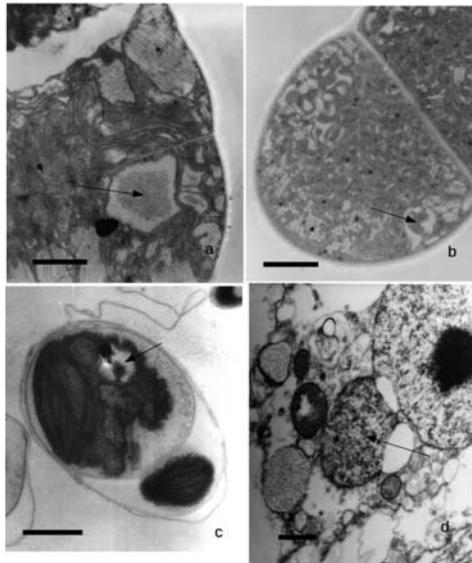


Fig. 1

1a: *Arthrospira platensis*. A large area containing granular material at low electron density is surrounded by membrane (arrow). Also the gas vesicles (asterisk) appear to be surrounded by a limiting membrane. Bar = 800 nm. 1b: *A. platensis*. A portion of cytoplasm with the same electron density of the rest of the cytoplasm is surrounded by thylakoidal membranes (arrow). Some gas vesicles appear fragmented (asterisk). Bar = 2 μ m. 1c: *Dichtiosphaerium* cfr. *Ehrenbergianum*. A vacuole (arrow) is enveloping a portion of cytoplasm and a lipid droplet is forming in the same organelle. Bar = 800 nm. 1d: *Heterosigma hakashiwo*. A portion (arrow) is detaching (blebbing) from the main part of the nucleus. Bar = 200 nm.

The eukaryotic algae do not represent a monophyletic group, since some clades of algae derived from primary endosymbiosis and others from secondary endosymbiosis. For this reason, the cell machinery does not work the same in every group. Concordantly, also autophagy showed quite variable features from the archaeplastida group (red algae, Glaucophyta, green algae and land plants) with respect, for instance, to the heterokont algae. Some example are presented.

Some doubt arises also about the origin of autophagy, since this phenomenon is typical of eukaryotes and currently no evidence of sequence similarity to autophagy-related eukaryotic genes has been found in prokaryotes genomes. As a matter of fact, autophagy consists in an activity of intracellular membranes and, by definition, prokaryotes should not have autonomous membranes with respect to the plasmamembrane. Nevertheless, cyanobacteria have a very developed membrane system, mainly devoted to photosynthesis, but also to the formation of gas vesicles. In cyanobacteria some open space, apparently surrounded by membranes, and containing granular material can be observed (Dvorak et al. 2015, Capelli et al. 2017), while there is evidence of protein recycling, particularly Rubisco. In *Arthrospira platensis* Gomont, some cytoplasm space containing granular material appear to get surrounded apparently by thylakoids and then to reduce their electron density (Fig. 1a and 1b). This observation may be a first hint to explain the strange autophagic activity performed by plastids in some plants (van Doorn, Papini 2013, Papini et al. 2014).

While in green algae the ultrastructural features are relatively similar to autophagy in land plants as in *Dichtiosphaerium* cfr. *ehrenbergianum* Nägeli (Fig. 1c), in the more distantly related *Heterosigma hakashiwo* (Hada) Hada ex Hara & Chihara (Raphidophyceae), belonging to one of the secondary endosymbiosis clades, autophagy assumes more extended features in the cytoplasm, particularly in case of starvation (Papini et al. 2017) and shows even evidence of nuclear blebbing (Fig. 1d), that is a typical feature of programmed cell death in metazoans.

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Speciation by time in diatoms: the case of *Pseudo-nitzschia allochroa* sp. nov., a cryptic species in the *P. delicatissima*-complex

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A new non-toxic species of the marine diatom genus *Pseudo-nitzschia*, *P. allochroa*, was isolated and cultivated from the Gulf of Naples and the Ionian Sea. Molecular differences in four phylogenetic markers (18S rDNA, LSU rDNA, ITS, *rbcl*) as well as the ITS2 secondary structure indicated this species to be distinctly separated from other congeneric species in the *P. delicatissima*-complex, some of which, i.e. *P. arenysensis*, *P. dolorosa* and *P. delicatissima* sensu stricto (s.s.), are very closely related to *P. allochroa* and occur in sympatry with it. The shape and ultrastructure of *P. allochroa* did not show any perceivable differences from those of *P. arenysensis* and *P. delicatissima*. Mating experiments showed sexual separation from the most closely related, *P. arenysensis*, definitely supporting the establishment of a new species. More than 60 strains of *P. allochroa* were isolated over the years from the LTER-MC station in the GON, and they were invariably recovered from the beginning of the summer throughout autumn. In contrast, *P. arenysensis*, *P. delicatissima* s.s. and *P. dolorosa* were recurrently found during the early spring annual peak. This clear separation over the annual cycle, also confirmed by metabarcoding data, may imply ecophysiological and functional differences between *P. allochroa* and the congeneric species. These results point at a possible mechanisms of speciation by time, whereby blooms of sympatric populations in different periods of the year may bring about sexual isolation and give rise to new taxa. As commonly accepted in the rest of the vegetal and animal realms, phenological patterns in diatoms are probably the result of genetic circannual clock modulated by environmental factors. Therefore, phenological signatures could be considered among phenotypic characters to be used in species delimitation.

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Effect of CO₂ supply on the growth and nutrient uptake in microalgae cultures for industrial applications

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Atmospheric CO₂ concentration has been rising since the Industrial Revolution and is predicted to double by the end of this century (IPCC 2014). Microalgae play an important role in the carbon cycle of marine ecosystems and their responses to elevated atmospheric CO₂ are of considerable interest (Li et al. 2013). The effects of CO₂ enrichment are variable and may depend on the availability of nutrients, since CO₂ supply leads to a higher growth rate and to an increased demand for nutrients.

CO₂ released by industrial activities contributes substantially to increase the atmospheric CO₂ level; thus it is becoming more and more important to find alternatives to control its release in the atmosphere (Beneman, Hughes 1997). Flue gases produced by processes such as anaerobic digestion, for instance, could be potentially used for microalgae culturing (de Godos et al. 2014), thus preventing their release and allowing the transformation of polluting gas fluxes into new and valuable microalgae products.

In this study physiological responses to CO₂ supply in two microalgal species that have potential industrial applications were investigated. In particular *Desmodesmus communis*, a green alga widely used for phytoremediation due to its high resistance to chemical and environmental stress, and the diatom *Phaeodactylum tricornerutum*, a PUFAs producer widely used in aquaculture, were studied using 1 L batch cultures (Fig. 1)



Fig. 2
Desmodesmus communis cultivation in 70 L photobioreactors.

or 70 L photobioreactors (Fig. 2, 3). Flue gases obtained by anaerobic digestion and enriched in CO₂ by the removal of the biogas fraction were also tested for microalgae culturing.

Results showed a marked increase of the growth and productivity in cultures with CO₂ addition respect to the ones grown with only air.

Moreover, a different use of macronutrients by algal cells and changes in cellular composition were observed when CO₂ was supplied. Preliminary experiments performed using flue gases supplied to *P. tricornerutum* cultures reported no inhibition of the algal growth and promising results at testing their potential use to cultivate microalgae.



Fig. 1
Batch cultures (1 L) of the diatom *Phaeodactylum tricornerutum*.



Fig. 3
CO₂ insufflation in a 70 L photobioreactor used for *Desmodesmus communis* cultivation.

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The ecology of the genus *Leptocylindrus* in a highly structured ecosystem, the northern Adriatic

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The marine diatom genus *Leptocylindrus* is a major component of phytoplankton blooms in coastal ecosystems worldwide; however, little is known about the ecology and meticulous species composition of this genus in the Adriatic Sea. Although *Leptocylindrus danicus* and *L. minimus* have been reported from numerous studies conducted in Adriatic, there has been neither unequivocal species identification nor focused examination of the temporal abundance of *Leptocylindrus* in this region. In a recent reappraisal of *Leptocylindrus* by Nanjappa et al. (2013) five species and one new genus were identified using both morphological and molecular markers. We here describe the temporal distribution of the genus *Leptocylindrus* based on 45 years of records, revealing that this diatom is a key component of the seasonal phytoplankton cycle, with greatest abundance in the spring and autumn. Using light and scanning electron microscopy as well as genetic analysis based on the nuclear-encoded rDNA regions, our study unambiguously revealed three species new for the Adriatic Sea, *Leptocylindrus hargravesii*, *L. convexus* and *L. aporus*. Furthermore, we investigated the growth, auxospore and resting spore formation of one of the *Leptocylindrus* species under different nutrient regimes. The ecology, physiology and evolution of this significant diatom genus should be further investigated.

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Canopy-forming algae and biodiversity: the structure and large-scale spatial variability of macroalgal assemblages associated to *Cystoseira* beds

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In the Mediterranean Sea, *Cystoseira* species are the most important canopy-forming algae in shallow rocky bottoms. The aim of the present study is to contribute to the knowledge of the structure of the *Cystoseira*-dominated assemblages through the assessment of a large-scale study. Specifically, we tested the hypothesis that the number of taxa, the abundance and the structure of epiphytic and epilithic macroalgae associated to the *Cystoseira* beds changed among different spatial scales. A hierarchical sampling design in a total of five sites across the Mediterranean Sea was used. A total of 118 taxa (species and genus level) associated to *Cystoseira* beds were identified. The Rhodophyta *Polysiphonia subulifera* and *Haliptilon virgatum* and the Ochrophyta *Sphacelaria cirrosa* and *Dictyota* spp were the most common epiphytic macroalgal taxa. The Rhodophyta *Laurencia obtusa*, and the Ochrophyta *Padina pavonica* and *Halopteris scoparia* were among the most abundant no-epiphytic macroalgae. The occurrence of small and large scale variability was highlighted, but the different assemblages composing the whole macro community associated to *Cystoseira* showed different patterns of spatial variability. The relative importance of the different scales of spatial variability highlighted can be of practical use to optimize sampling designs focused on the ecological problems of this habitat and monitoring plans.

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Phytoplankton stable carbon isotopes as a tool to monitor CO₂ leakage at Carbon Capture and Storage sites

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Carbon capture and storage technology (CCS) is expected to play a key role among strategies for the mitigation of climate change by reducing CO₂ emissions into atmosphere from fossil fuel combustion. Although a well-engineered storage site is not expected to leak, the risk of failure of CO₂ containment and the subsequent environmental impact is a major issue for the acceptance of this approach. This study aims to the evaluation of stable carbon isotope analysis as a tool for effective early warning of CO₂ migration from CCS, since different carbon sources have specific $\delta^{13}\text{C}$ (usually lower values in anthropogenic emissions than in natural CO₂).

Two culture experiments were conducted within 2L-photobioreactors under controlled conditions (light, temperature). In each experiment, the diatom *Thalassiosira rotula* was grown in two different media: one prepared with natural seawater and the other with artificial seawater, whose carbonate system was derived from anthropogenic CO₂. Both media were equally supplemented with nutrients (silicate, nitrate and phosphate), metals and vitamins. Daily variations of $\delta^{13}\text{C}$ in dissolved inorganic carbon (DIC) and in diatoms were analysed in order to study the phytoplankton response.

According to the experiments results, differences exist in microalgae isotope composition between natural and anthropogenic conditions. In natural medium, the diatoms $\delta^{13}\text{C}$ values did not show important deviations from the starting value ($-24.4 \pm 0.3\text{‰}$), whereas, in the anthropogenic CO₂ condition, the uptake of ¹³C-depleted DIC ($-44.7 \pm 0.8\text{‰}$) resulted in the decreasing values of microalgae carbon isotope composition (until -44.4‰). The rapid change in phytoplankton isotope composition confirms the method tested as reliable for identifying different CO₂ sources.

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Phytoplankton traits, functional groups and community organization

L. Roselli, E. Litchman

A central goal in ecology is to understand and predict the effects of environmental change on ecological communities and the consequences of community change for ecosystem functioning. Searching for common patterns and decoding underlying mechanisms from those patterns is how scientists can contribute to sustainable ecosystem management and conservation. Trait-based approaches have become increasingly successful in community ecology in general and phytoplankton communities in particular. Phytoplankton communities have a rich history as model systems in community ecology and are ideally suited for applying and further developing trait-based concepts. Understanding trait changes is key to better forecasting community responses to environmental impacts, including anthropogenic global change. In the context of maintenance of biodiversity and ecological functions, microbial ecologists face the challenge of linking individual level variability in functional traits to larger scale ecosystem processes. Phytoplankton cell size and shape are key traits under selection by environmental filters and species interactions. Here, we explore how size and shape enter the diversity game. How does taxonomic and morpho-functional community structure vary at different spatial scales? What are the potential drivers shaping the structure of phytoplankton communities? We explore these questions by looking at the individual level variability in taxonomic and morphological traits in a biogeographical snapshot of natural phytoplankton communities in coastal ecosystems around the globe. Species diversity was more variable than trait diversity from local to global spatial scales. We suggest that structural organization of phytoplankton communities in coastal ecosystems may follow a hierarchical pattern of trait organization, where a different combination of multiple functional traits may represent effective strategies and promote success under given environmental conditions as a resolution of Hutchinson's paradox.

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Un approccio integrato per la rimozione di fosforo e metalli pesanti accoppiata alla produzione di lipidi ad opera della microalga verde *Desmodesmus* sp.

L. Rugnini, G. Costa, R. Congestri, S. Antonaroli, L. Bruno

La qualità dei corpi idrici superficiali è fortemente influenzata dalle attività antropiche che causano emissioni diffuse e / o puntiformi di inquinanti organici e inorganici. Gli effluenti ricchi di nutrienti come l'azoto (N) e il fosforo (P) possono dar luogo a fenomeni come l'eutrofizzazione (Schindler et al. 2008), mentre la presenza di metalli può rappresentare un serio rischio sia per la salute umana che per l'ecosistema (Torres et al. 2017). Le microalghie sono in grado di sottrarre i nutrienti in eccesso nelle acque reflue e di resistere alla presenza di metalli tossici accumulandoli all'interno delle cellule, caratteristica che ne permette l'utilizzo nel settore del biorimedio (Gismondi et al. 2016, Rugnini et al., 2017, 2018). L'impiego di microalghie per il trattamento delle acque reflue permette, inoltre, di ottenere una biomassa che può essere, ad esempio, impiegata per la produzione di biocarburanti come biodiesel, etanolo, bioidrogeno (Bruno et al. 2012, Bux, Chisti 2016, Gismondi et al. 2016). In questo lavoro è stata valutata la possibilità di utilizzare la microalga verde *Desmodesmus* sp. (Fig. 1) sia per il biorimedio delle acque reflue

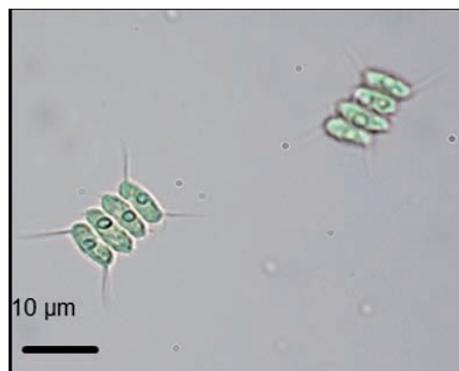


Fig. 1
Immagine al microscopio ottico di cenobi di *Desmodesmus* sp.

dall'eccesso di P e metalli pesanti quali rame (Cu) e nichel (Ni), sia per la produzione di biodiesel.

Sono stati effettuati due esperimenti di crescita in coltura, indicati come *Run 1* e *Run 2*, in fotobioreattori a basso costo della capacità 10 L (Fig. 2) utilizzando come mezzo di coltura il terreno BG11 (Rippka et al. 1979) modificato per il contenuto di P che era pari a 4.55 mg L⁻¹ per simulare le concentrazioni osservate in effluenti di impianti di depurazione civili o industriali (Water Environment Federation 2010, Cai et al. 2013). Inoltre, nel *Run 2*, la coltura di *Desmodesmus* sp. è stata anche esposta a una soluzione bimetallica di Cu e Ni e dopo 2 giorni di esposizione/contatto è stata valutata la capacità di biosorbimento delle cellule. La biomassa ottenuta in entrambi i *Run* è stata sottoposta ad analisi della capacità di rimuovere il P e ad estrazione dei lipidi, che successivamente sono stati analizzati mediante gas-cromatografia per la caratterizzazione degli esteri metilici degli acidi grassi (FAME).

I risultati hanno dimostrato che *Desmodesmus* sp. è in grado di rimuovere tra il 96 e il 100% del P presente, impiegando 21 giorni nel *Run 1* e 14 giorni nel *Run 2*, mentre sono stati sufficienti solo 2 giorni di contatto microalga-metallo per la rimozione del 94% del Cu totale e l'85% di Ni (Rugnini et al. 2018). Molti studi (Kumar et al. 2015, Rugnini et al. 2017) riportano che in soluzioni multi-metalliche che contengono Cu e Ni l'affinità per il Cu è sempre maggiore a quella del Ni, come ottenuto anche in questo studio, probabilmente a causa di una più forte interazione tra ione metallico e gruppi carbossilati della parete cellulare delle microalghie. È noto come il contenuto lipidico delle alghe dipenda da diversi fattori, come la disponibilità dei nutrienti nel mezzo di coltura, la salinità, l'intensità luminosa, e la presenza di metalli e altri contaminanti che possono indurre stress ossidativi causa di accumulo di lipidi come meccanismo di difesa (Torres et al. 2017). In questo studio, l'estrazione e lo studio dei profili lipidici hanno rivelato che la presenza di metalli ha diminuito la resa lipidica, ma, secondo gli standard europei (EN 14214) e statu-

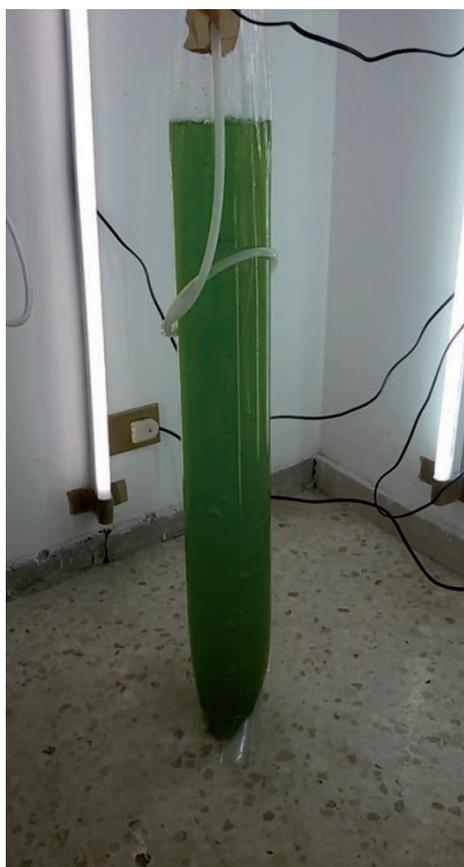


Fig. 2
Fotobioreattore da 10 L costituito da buste di polietilene impiegato per gli esperimenti indicati come *Run 1* e *Run 2*.

nitensi (ASTM-D6751), la qualità del biodiesel ottenuto dalla biomassa esposta ai metalli (*Run 2*) era superiore a quella del biodiesel ottenuto dalla biomassa non esposta (*Run 1*), poiché risultava maggiore la quantità di lipidi saturi e monoinsaturi accumulati.

Questo studio ha dimostrato, quindi, l'efficacia dell'utilizzo della microalga verde *Desmodesmus* sp. per la rimozione di contaminanti come P, Cu e Ni. Tale applicazione può essere integrata con successo all'uso della biomassa ottenuta come materia prima per la produzione di biocarburanti, offrendo una possibile soluzione alternativa a più problemi ambientali.

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The use of Marine Resources for Restoration and Conservation of Artworks

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In the conservation of artifacts the use of natural materials has been increasing in recent year, especially at Vatican Museums, thanks to the work of the Scientific Laboratories. In many cases, natural materials are progressively replacing synthetic ones, allowing less invasive interventions on artworks and a minor impact on operators and the Environment. The emergence of these new restoration methodologies represents, thus, a real change in approaches, leading restorers towards the principle of sustainability in the practice of conservation. In the restoration Laboratories of Vatican Museums, chemists, naturalists, biologists, and restorers work together to find new restoration solutions through the use of natural materials. In this context, marine resources represent an extraordinary source of innovative products to be tested in the field of Cultural Heritage. Among the natural materials for consolidation, "Funori", a polysaccharide extracted from the red alga *Gloiopeltis furcata* (Postels & Ruprecht) J. Agardh of the Pacific Ocean is now commonly used. The matrix from which the Funori is obtained is soluble in hot water and consists of a mixture of polysaccharides, sulphates, lipids, proteins, salts, and dyes (Izumi 1971). *G. furcata* is purified in laboratory and its extract is used by restorers on paper, wood, metal, and wall paintings (Michel 2011). At the same time, studies and experiments on the red Mediterranean alga *Sphaerococcus coronopifolius* Stackhouse, are ongoing at Vatican Museums. Several tests have shown that the extract obtained from *S. coronopifolius* has properties similar to those of Funori. Moreover, it seems to may assure a better yield in terms of pH, conductivity and adhesiveness (Fratini et al 2016). Recent studies have demonstrated that the species reproduces on the Lazio coasts through a direct-type of life history, ensuring a less variability in the composition and yield of the natural products synthesized by it (Abdelahad et al 2016). Finally, the use of *Posidonia oceanica* (L.) Delile for restoration purposes is to be also remarked (Lomoro et al 2011). The egocropiles of this marine plant are treated in laboratory and used as stucco for woodwork. Tests and applications on *P. oceanica* are still ongoing at the Scientific Laboratories (Minnini et al. 2009, Parente et al. 2009).

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Acclimatazione all'acidificazione in *Caulerpa cylindracea* Sonder

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Il continuo aumento di gas serra, in particolare di anidride carbonica (CO₂) rilasciata nell'atmosfera dalle crescenti attività antropiche attraverso l'uso di combustibili fossili, è causa di importanti cambiamenti climatici i cui effetti più eclatanti si manifestano in un aumento della temperatura e in una devastante siccità (IPCC, 2014). Nell'ambiente marino il riscaldamento si sovrappone ad un altro effetto, l'acidificazione degli oceani, dal momento che un terzo dell'anidride carbonica emessa nell'atmosfera viene assorbita dalle grandi masse d'acqua che ricoprono il nostro pianeta (Sabine et al. 2004).

Mentre i primi studi sugli effetti dell'acidificazione sono stati condotti per lo più in laboratorio o in mesocosmi semi-naturali utilizzando organismi con alti tassi di crescita e brevi cicli vitali (Kim et al. 2016), negli ultimi dieci anni l'attenzione si è spostata verso quei siti marini naturalmente acidificati dalla presenza di emissioni gassose di varia natura e che perciò possono rappresentare laboratori naturali in cui validare la coerenza dei risultati ottenuti in sistemi artificiali (Fabry et al. 2008, Koch et al. 2013, Kroeker et al. 2013).

L'obiettivo di questa ricerca è stato quello di studiare le risposte di *Caulerpa cylindracea* Sonder a variazioni di pH. Questa specie invasiva di origine tropicale, grazie alla sua elevata plasticità, è in grado di colonizzare vari habitat con condizioni di luce e temperatura estremamente varie (Raniello et al. 2006, Flagella et al. 2008).

Per valutare la risposta della macrofita all'acidificazione, sono stati condotti *in situ* trapianti incrociati di talli tra aree a pH 8.01 e aree a pH 7.50, nel sito del Castello Aragonese di Ischia (Napoli), dove emissioni naturali di CO₂ creano un gradiente di acidità compreso tra 8.1 e 6.7 (Hall-Spencer et al. 2008, Porzio et al. 2011, 2013). Gli effetti legati alle variazioni di pH ambientale sulla specie in esame sono stati studiati al microscopio ottico ed elettronico a trasmissione. È stata inoltre valutata l'efficienza fotosintetica attraverso variazioni dell'attività fotochimica (Diving PAM) e del contenuto in pigmenti. Sono state analizzate le risposte ai cambiamenti di pH a breve termine (3, 7, 14 giorni dal trapianto) e la capacità di recupero dopo il riposizionamento dei talli alle condizioni di origine, per valutare l'eventuale reversibilità delle modificazioni subite. Inoltre, il confronto tra le due popolazioni naturali a pH 7,50 e a pH 8,01 ha permesso di valutare gli effetti su lunga scala temporale.

Mentre i risultati relativi agli effetti su lunga scala temporale non hanno evidenziato importanti alterazioni fisiologiche e ultrastrutturali, nel breve periodo sono state riscontrate modificazioni significative. I talli trapiantati da pH 7.50 a 8.01, infatti, hanno risposto positivamente all'aumento del pH, mostrando un aumento progressivo del numero di cloroplasti all'aumentare del tempo di esposizione alla nuova condizione ambientale, un aumento del rapporto Chl a/Chl b e una diminuzione del contenuto in carotenoidi dal 3° al 14° giorno dal trapianto. Inoltre, è stato riscontrato un miglioramento dell'efficienza fotosintetica ed una diminuzione della dissipazione termica dell'energia di eccitazione, segnali questi di un passaggio verso un ambiente favorevole.

Al contrario, il trasferimento da pH 8.01 a pH 7.50 ha innescato risposte negative evidenziate dalla diminuzione nel numero di cloroplasti, dall'aumento delle dilatazioni tra le membrane tilacoidali e da importanti accumuli di amido. Queste variazioni ultrastrutturali sono state confermate dalla diminuita efficienza fotosintetica, dall'aumento dei carotenoidi e del rendimento non fotochimico e dalla diminuzione delle clorofille, sottolineando uno stress per la macroalga.

D'altra parte, il fatto che *C. cylindracea* sia presente naturalmente in siti a pH 8,01 e a pH 7,05, ci porta a supporre che le risposte allo stress da acidificazione messe in atto nel breve periodo siano transitorie e che tale specie potrebbe acclimatarsi al progressivo acidificarsi dei mari, nello scenario dei cambiamenti climatici in atto.

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Intensive cultivation of *Staurosirella pinnata* (Bacillariophyceae) for mycosporine-like aminoacid characterization and bioactivity on melanoma cell line

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Mycosporine-like amino acids (MAAs) are a class of small, water-soluble molecules known for their capability of absorbing UV radiation and widespread across a variety of algae and cyanobacteria. Some red algae MAAs showed bioactivity with potential in drug discovery, but no evidence yet exist on MAA production, composition and potential application in cultured diatoms.

In this work, one brackish strain of *Staurosirella pinnata* (Ehrenberg) D.M. Williams & Round, VRUC 290, was mass cultivated in indoor polyethylene photobioreactors (10 L) to produce biomass for MAA extraction using methanol/water (20:80 v/v). Extracts were then characterized using LC-MS and subsequently analyzed for their anti-oxidant activity with 2,2-diphenyl-1-picrylhydrazyl (DPPH) assay and, finally, tested for their effect on melanoma cells (CHL-1) by flow cytometry.

Chromatographic profiles allowed identification of four different mycosporines, two of which mycosporine-glycine and euhalothece, present in relatively high amounts compared to concentrations recorded for other two diatom strains *Phaeodactylum tricornutum* and *Thalassiosira weissflogii* (SAG Culture Collection) cultivated in the same conditions in our laboratory. The radical scavenging activity of *S. pinnata* extract was assessed by DPPH assay showing high anti-oxidant activity, with an IC_{50} of $15.0 \pm 0.1 \mu\text{g}$.

Bioactivity of the extract was then tested on human melanoma CHL-1 cells and cytofluorimetric analyses (Fluorescence-Activated Cell Sorting) evidenced a strong cytotoxic effect of the after 24h and 48h treatments. Finally, alteration of cell cycle was observed after 24 and 48h treatments. CHL-1 cells showed an increase in cells in G1 and G2/M phases while a decrease of cells in S phase was measured. These results suggest an anti-proliferative effect of the *S. pinnata* extract on the melanoma cell line.

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Success of aquatic angiosperm transplantation in the northern Venice Lagoon

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The project SeResto (LIFE12 NAT/IT/000331) aimed at the recovering of the northern part of the Venice Lagoon by transplantation of aquatic angiosperms. The main anthropogenic impacts that affected the lagoon in the second part of the last century reducing aquatic angiosperm prairies, since the mid-2000s declined. The banning of phosphorus from the formulation of detergents (October 1988), the installation of urban waste water treatment plants in the hinterland and a general management of nutrient inputs in the lagoon, triggered a progressive reduction of the trophic conditions and blooms of Ulvaceae restoring conditions suitable for plant recruitment.

The objective was to trigger the natural re-colonization in a confined area where plant presence was negligible by mean of widespread, manual and low-cost transplantations of small sods or single rhizomes, mainly of *Zostera marina* and *Z. noltei*, avoiding any impact on the shallow bottoms. Aquatic angiosperms are environmental engineers which provide many environmental services and suitable conditions for benthic, fish and bird fauna. Their presence is the first condition for habitat recovering and the restoration of pristine or almost pristine environments.

After three years from the beginning of the project ca. 44700 rhizomes have been transplanted in 35 sites placed in an area of ca. 36 km². Transplantations have been successful in 32 sites with a mean coverage of 60% and many areas have been completely colonized. However, results were very different among sites and were negatively affected by raining conditions and water temperature whose extreme values can severely hamper their spread. Despite the different and unpredictable climatic conditions now aquatic angiosperms have colonized all the bottoms close to salt marshes and the canal edges. They are self-sustaining and expand exponentially because they produce a high number of seeds that colonize all the suitable areas. If extreme climatic changes are not taking place and other human impacts will be not occur, in a few years the environmental conditions of the northern lagoon will be restored as before the anthropogenic impacts.

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Investigating silver nanoparticles toxicity on marine macroalgae (*Ulva rigida*) using biochemical and physiological biomarkers

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Keywords: biomarkers, silver nanoparticles, *Ulva rigida*, Venice Lagoon

Despite the high ecological and economical importance of transitional environments and their high productivity, they represent a sink for many pollutants including metals. The steady increase in the production and release of products, containing silver nanoparticles (AgNP), could lead to an increasing dispersion and diffusion of silver into transitional environments. In this study, *in vitro* experiments were conducted to examine the toxicity of AgNP using the marine macroalga *Ulva rigida* C. Agardh, the most common species in transitional environments. The toxicity of both AgNP and aqueous silver (AgNO₃) was estimated based on the implementation of both biochemical (lipid peroxidation) and physiological (primary production) biomarkers with conditions similar to the Venice Lagoon environment.

The primary production showed an adverse effect of AgNP starting from 0.1 ppm reaching complete inhibition at 5ppm. Conversely, AgNO₃ displayed a severe inhibition of primary production in *U. rigida* already at 0.01 ppm.

The lipid peroxidation, measured by malondialdehyde levels, showed that the lowest observed stress effect was at 1 ppm of AgNP. The oxidative stress generated by AgNP steadily increased up to a concentration of 30 ppm resulting in membrane disruption. In contrast, the oxidative stress generated by AgNO₃ was less significant, proving that AgNO₃ was very toxic (as measured by primary production) but did not produce stress at cellular level. The silver bioaccumulated by *U. rigida* from an AgNP solution of 1 ppm reached 8.5 mg Kg⁻¹fw and the bioaccumulation reached a saturation point at 11.3 mg Kg⁻¹fw from a 5ppm AgNP solution.

The concentration of silver in the water column don't exceed the ppt range, which is not toxic for *U. rigida*, but the reported concentrations for benthic marine organisms from the Venice Lagoon were in the ppm range. These concentrations of AgNP were found to be high enough when bioaccumulated to elicit a phytotoxic effect.

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Acidification and temperature effects on the gene expression of the coccolithophore *Pleurochrysis pseudoroscoffensis*

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Coccolithophores are nanoplanktonic microalgae characterized by an exo-skeleton composed of minute calcified plates and play an important role in control of climate changes because they are involved in the sulfur cycle for the production of dimethylsulfide as well as in the carbon cycle for the photosynthesis and calcification processes. Recent studies on the ocean acidification due to the carbon dioxide increase in the atmosphere have provided evidence on a possible impact on coccolithophore calcification.

The aim of this study was: a) to investigate possible effects on gene expression of the coccolithophore *Pleurochrysis pseudoroscoffensis* due to changes in $p\text{CO}_2$ and temperature; b) to assemble de novo the genome and transcriptome of the coccolithophore.

In the framework of the project Acid.it (Costruzione di conoscenze e di strumenti a supporto della definizione di strategie di mitigazione ed adattamento agli effetti dell'acidificazione marina, con particolare riferimento ai mari italiani), culture experiments were performed in triplicate in 2.5 L-photobioreactors testing four $p\text{CO}_2$ /temperature conditions: control (21 °C and 400 ppm CO_2), high temperature (25 °C and 400 ppm CO_2), high $p\text{CO}_2$ (21 °C and 700 ppm CO_2) and high temperature and $p\text{CO}_2$ (25 °C and 700 ppm CO_2). Samples were collected at two different timing: after 11 and 20 days. The RNA from samples was isolated with a commercial kit, and the cDNA library were prepared for the sequencing with the Illumina HiSeq 3000/4000 technology. The reads obtained from the sequencing were trimmed and assembled with the Trinity software, the transcriptome is about 51000 contigs. The functional annotation was made with the software Trinotate, and for evaluate differentially expressed genes was used the program CLC Genomics Workbench.

Considering the gene expression, the only effect of CO_2 did not result in significant variations, whereas the only temperature had a more significant effect. However, these two factors seemed to show a combined effect, with a strong gene up-regulation and down-regulation at the end of the culture. The genes down regulated were related to protein transcription and translation, so to presume a complete stop of the protein synthesis. Among the upregulated genes, there were some involved in several metabolic pathways mainly related to nitrogen synthesis, but also genes involved in the coccolithogenesis and a carbonic anhydrase, suggesting a possible alteration of the production of the calcium carbonate skeleton.

Instead, for the genome we isolated the DNA and sent it to the sequencer center, where they prepare the library and sequenced them with the Illumina HiSeq 3000/4000 technology. The reads were then assembled with four different software (CLC G.W., Soap, Spades2 and Abyss). The better output, based on different parameters was the one with Abyss, and with another program (Jellyfish) we were able to estimate the genome size which is about 213Mb.

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The Ecology of One Cosmopolitan, One Newly Introduced and One Occasionally Advected Species from the Genus *Skeletonema* in a Highly Structured Ecosystem, the Northern Adriatic

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The diatom genus *Skeletonema* is globally distributed and often an important constituent of the phytoplankton community. In the marine phytoplankton of the northern Adriatic Sea, we found three species of the genus *Skeletonema*: *Skeletonema menzelii*, *S. marinoi* and *S. grevillei*. Making use of the steep ecological gradients that characterize the northern Adriatic, along which we could observe those species, we report here on the ecological circumstances under which those species thrive and how their respective populations are globally connected. This is the first detailed ecological study for the species *S. grevillei*. This study is also the first report for *S. grevillei* for the Adriatic Sea and Mediterranean, together with additional electron microscopic details on fresh in situ samples for this species. *S. marinoi* appears to clearly prefer strong freshwater influence and high nutrient concentrations delivered by low salinity waters. It can outcompete other diatom species and dominate microphytoplankton blooms. *S. grevillei* on the other hand appears to thrive in high nutrient concentrations triggered by water column mixing. It also appears to prefer higher salinity waters and coastal embayments. Genetic analysis of *S. grevillei* demonstrated a peculiar dissimilarity with isolates from coastal waters off Yemen, India, Oman and China. However, a closely related sequence was isolated from coastal waters off Japan. These results indicate that *S. grevillei* is an introduced species, possibly transported by ballast waters. *S. menzelii* is a sporadic visitor in the northern Adriatic, advected from rather oligotrophic middle Adriatic waters and never dominates the phytoplankton community in the northern Adriatic.

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Preliminary study of primary production and nutrient uptake by two seagrasses (*Zostera marina* and *Zostera noltei*) in the Venice Lagoon

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Seagrass meadows are among the most productive ecosystems on Earth, that provide high-value ecosystem service and contribute to carbon sequestration. Nutrient and light availability are the primary physical factors controlling seagrass growth. The median carbon, nitrogen and phosphorus contents in seagrass leaves translate into a median atomic C:N:P ratio of 432:20:1. If we compare these values with the Redfield ratio for the production of organic matter by phytoplankton (C:N:P 106:16:1), we can deduce that seagrasses need about a quartet of the nitrogen and the phosphorus that are necessary for phytoplankton.

This is the first study *in situ* of seagrass primary production and nutrient uptake from the water column in the Venice Lagoon. We tested the Net Community Production (NCP) and the nutrient uptake (nitrogen and phosphorus) of *Zostera marina* and *Z. noltei* through benthic incubation chambers. This work was carried out in summer (July and August 2017). The result showed that the NCP of *Z. marina* decreased from July (mean 0.108 ± 0.02 mmol m⁻² h⁻¹ g(ftw)⁻¹) to August (mean 0.053 ± 0.08 mmol m⁻² h⁻¹ g(ftw)⁻¹), according to the growth rates of this specie. Moreover, the comparison of *Z. marina* and *Z. noltei* NCP, during the August sampling, highlighted that the species with the highest NCP was *Z. noltei*, with a mean of 0.709 ± 0.35 mmol m⁻² h⁻¹ g(ftw)⁻¹ against a mean of 0.108 ± 0.02 mmol m⁻² h⁻¹ g(ftw)⁻¹ of *Z. marina*. Indeed, in this month, the growth rates of *Z. noltei* increased while that of *Z. marina* decreased.

Finally, this study displayed that the nutrient uptake for both the species was higher for ammonium, with a mean of 17.1 nM h⁻¹ g(ftw)⁻¹ for *Z. marina* and a mean of 23.5 nM h⁻¹ g(ftw)⁻¹ for *Z. noltei*, while the phosphorus and nitrate uptake were lower. In particular, the mean for both species of phosphorus was 5.2 nM h⁻¹ g(ftw)⁻¹ and of nitrate was 1.1 nM h⁻¹ g(ftw)⁻¹.

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Application of taxonomic and morpho-functional properties of phytoplankton communities to water quality assessment for artificial lakes in the Mediterranean Ecoregion

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Starting in September 2010, a monitoring program was developed for evaluating the water quality in six artificial lakes (reservoirs) of the Apulia Region (Southern Italy). Lake ecological status was evaluated according to the requirements of the 2000/60/EC Water Framework Directive. Phytoplankton taxonomic composition, biovolume and chlorophyll a data were used in an application of the Italian methods (Italian Phytoplankton Assessment Method or New Italian Method) for water quality assessment of artificial lakes. Even though all the reservoirs were classified as being in good ecological status, the phytoplankton communities of the six reservoirs were different and characterized by highly site-specific components and by distinctive dominant taxa. The influences of environmental variables on the phytoplankton communities were analyzed. Results of Constrained Coordinate Analysis (CCA) indicated strong influences of the trophic gradient and water depth on the phytoplankton assemblages, but significant factors are also the surrounding landscape and the origin and the history of the lakes. A functional traits-based analysis was carried out to support the water quality assessment tools. Twenty-one morpho-functional groups were identified. Most of them were common in all six lakes, so functional properties clearly recur among different phytoplankton species.

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