



Drivers of Cyanobacterial Blooms in a Hypertrophic Lagoon

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Bartoli M, Zilius M, Bresciani M, Vaiciute D, Vybernaite-Lubiene I, Petkuviene J, Giordani G, Daunys D, Ruginis T, Benelli S, Giardino C, Bukaveckas PA, Zemlys P, Griniene E, Gasiunaite ZR, Lesutiene J, Pilkaitytė R and Baziukas-Razinkovas A (2018) Drivers of Cyanobacterial Blooms in a Hypertrophic Lagoon. Front. Mar. Sci. 5:434. doi: 10.3389/fmars.2018.00434 The Curonian Lagoon is Europe's largest lagoon and one of the most seriously impacted by harmful blooms of cyanobacteria. Intensive studies over the past 20 years have allowed us to identify the major drivers determining the composition and spatial extent of hyperblooms in this system. We summarize and discuss the main outcomes of these studies and provide an updated, conceptual scheme of the multiple interactions between climatic and hydrologic factors, and their influence on internal and external processes that promote cyanobacterial blooms. Retrospective analysis of remote sensed images demonstrated the variability of blooms in terms of timing, extension and intensity, suggesting that they occur only under specific circumstances. Monthly analysis of nutrient loads and stoichiometry from the principal tributary (Nemunas River) revealed large interannual differences in the delivery of key elements, but summer months were always characterized by a strong dissolved inorganic N (and Si) limitation, that depresses diatoms and favors the dominance of cyanobacteria. Cyanobacteria blooms occurred during high water temperatures, long water residence time and low-wind conditions. The blooms induce transient (night-time) hypoxia, which stimulates the release of iron-bound P, producing a positive feedback for blooms of N-fixing cyanobacteria. Consumermediated nutrient recycling by dreissenid mussels, chironomid larvae, cyprinids and large bird colonies, may also affect P availability, but their role as drivers of cyanobacteria blooms is understudied.

Keywords: Curonian Lagoon, nitrogen, phosphorus, silica, fluxes, stoichiometry, remote sensing, cyanobacteria

INTRODUCTION

Human activities impact biogeochemical cycles, biological communities and ecosystem functioning of inland and coastal waters on a global scale (Bernot and Dodds, 2005; Muhlolland et al., 2008; Paerl, 2009; Han and Allan, 2012). Estuaries and lagoons have become enriched with nutrients due to wastewater discharge, aquaculture, and agriculture (Galloway et al., 2008; Paerl, 2009). Excess nutrients result in blooms, where algal biomass accumulates and exceeds the mineralization capacity of the heterotrophic community (Valiela et al., 1997). In fresh-brackish waters, algal

blooms may include one or more types of harmful cyanobacteria, resulting in the presence of cyanotoxins. The development of hypoxic or anoxic conditions can lead to die-offs of fish and benthic organisms (Norkko and Bonsdorff, 1996; Ye et al., 2011). In addition, changes in food web structure brought about by invasive species may accelerate eutrophication by reducing grazing pressure and allowing the proliferation of algae, including toxic forms (Carpenter et al., 1998; Rabalais et al., 2002). The alteration of nutrient stoichiometry (sensu Redfield) and changes in climate (e.g., warmer temperatures, precipitation timing and intensity) have also received attention as potential drivers of harmful algal blooms (Cloern, 2001; Yunev et al., 2007; Moore et al., 2008; Howarth et al., 2011). In this review, we analyze the drivers of algal blooms in the Curonian Lagoon, a hypereutrophic freshwater estuary. We discuss the relevance of nutrient loads and their stoichiometry on algal blooms, we analyze how algal blooms affect the ecosystem functioning (e.g., nutrient mass balances) and provide a mechanistic interpretation for positive feedbacks promoting the dominance of cyanobacteria.

THE CURONIAN LAGOON: GENERAL FEATURES OF A HYPERTROPHIC FRESHWATER ESTUARY

The Curonian Lagoon is a large (surface area = 1500 km^2), shallow (mean depth = 3.5 m) waterbody located along the south-eastern portion of the Baltic Sea (Figure 1). The Curonian Spit (a UNESCO heritage site) divides the lagoon from the Baltic Sea. The main source of water and nutrients is the Nemunas River, although the lagoon also exchanges water with the Baltic Sea via the narrow Klaipeda Strait (Vybernaite-Lubiene et al., 2017). Exchange of water through the strait is episodic; during wind-driven forcing events, the salinity of the lagoon rises to \sim 7. The principal tributary (Nemunas River) bisects the lagoon such that the northern lagoon is subject to greater fluvial (and marine) influence, whereas the southern portion of the lagoon is more lentic, and has a longer water residence time (Umgiesser et al., 2016). The lagoon has a relatively small hydrologic loading factor (ratio of watershed area to surface area), which makes this system similar to a flowthrough reactor, and provides an opportunity for mass balance studies (Bresciani et al., 2012; Zilius et al., 2014; Vybernaite-Lubiene et al., 2017). Prior work by our multidisciplinary and multinational team has included assessment of nutrient loads from the Nemunas watershed, application of hydraulic models to simulate water circulation in response to changing discharge and wind conditions, investigation of seasonal dynamics of biogeochemical cycles, and use of satellite remote sensing to monitor phytoplankton blooms (Vaičiūtė et al., 2015; Petkuviene et al., 2016; Umgiesser et al., 2016; Vybernaite-Lubiene et al., 2017, 2018). Our work at this site has also benefitted from longterm monitoring carried out by the Marine Research Department of the Lithuanian Ministry of Environment.

Data arising from these efforts have helped to guide management of the Curonian Lagoon via a number of national and international programs (HELCOM Baltic Sea Action Plan, European Water Framework Directive, various habitat and bird conservation initiatives). Despite the intensive studies carried out in the lagoon, there remains the question whether and how cyanobacteria blooms can be mitigated. These blooms extend over large areas of the lagoon and negatively impact ecosystem functions, including tourism and recreational activities, as well as local fisheries (Giardino et al., 2010; Belykh et al., 2013; Šulčius et al., 2015). The use of science for informing management decisions is dependent upon the interpretation and integration of available data, which is the focus of this paper.

SEASONAL SUCCESSION OF PHYTOPLANKTON

Detailed studies of plankton communities in the Curonian Lagoon have examined seasonal patterns, species interactions, production of cyanotoxins and the role of phytoplankton in food web energetics (e.g., Pilkaitytė and Razinkovas, 2006, Razinkovas, 2007; Lesutienė et al., 2014; Bukaveckas et al., 2017). Diatoms dominate the spring phytoplankton community, after which, following a short clear-water phase, cyanobacteria biomass increases (Gasiunaitė et al., 2005; Pilkaitytė and Razinkovas, 2007). Fresh-brackish species dominate the phytoplankton community of the Curonian Lagoon. Stephanodiscus hantzschii, Diatoma tenuis, Aulacoseira islandica, Asterionella formosa are the dominant diatom species during spring while the N-fixing cyanobacteria Aphanizomenon flosaquae, Dolichospermum affine, D. flosaquae, as well as other cyanobacteria such as Microcystis aeruginosa, M. wesenbergii, M. viridis, and Planktothrix agardhii contribute to the summer biomass peak (Pilkaitytė and Razinkovas, 2007; Gasiunaitė et al., 2008). According to longterm monitoring data (2001-2012), monthly average chlorophyll a (chl-a) concentrations reach 47 \pm 14 mg m⁻³ during the spring diatom bloom and 96 \pm 56 mg m⁻³ during the summer bloom (Marine Research Department of the Lithuanian Ministry of Environment).

WIND EFFECTS ON ALGAL BLOOM DEVELOPMENT

Algal blooms in the Curonian Lagoon have been tracked since the 1930's via synoptic sampling (Schmidt-Ries, 1940). More recently, satellite remote sensing has substantially improved our ability to track the spatial and temporal dynamics of bloom events and draw links to local weather conditions. The first attempt to map algal blooms in the Curonian Lagoon utilized the MEdium Resolution Imaging Spectrometer (MERIS) on board the Envisat-1 satellite (Giardino et al., 2010; Bresciani et al., 2012; Vaičiūtė et al., 2015). The combination of high spatial resolution (300 m) and short revisit time (2–3 days) greatly enhanced our ability to map chl-a. More recently, the Operational Land Imager (OLI, on board Landsat-8) and Multispectral Instrument (MSI, on board Sentinel-2A/B) have further enhanced spatial resolution (10–30 m) and allowed us



to investigate the patchy distribution of cyanobacteria blooms (INFORM, 2017).

Results based on a large number of images from 2004 to 2016 revealed temporal variability and small-scale spatial patchiness of chl-a (Bresciani et al., 2012, 2014; Vaiči \overline{u} tė et al., 2015). The southern part of the lagoon exhibited high chl-a (up to 500 mg m⁻³) while the northern areas were characterized by lower values (~50 mg m⁻³) (Bresciani et al., 2012; **Figure 2**). Differences between the northern and southern portions of the lagoon were documented by earlier studies (Olenina, 1998; Krevs et al., 2007). However, the use of satellite images allowed us to identify hot spots of chl-a (up to 400 mg m⁻³) and the presence of surface scums (e.g., Bresciani et al., 2014; **Figure 2**). Highest concentrations were coincident with prevailing wind conditions, suggesting that wind speed and direction was a significant driver for spatial distribution of positively buoyant cyanobacteria (Bresciani et al., 2014).

Wind speed affects not only the spatial distribution of cyanobacteria, but also influences water column mixing. Wind speeds less than 2 m s⁻¹ are common and allow for the development of transient (daytime) thermal gradients within the water column (Zilius et al., 2014). The lagoon, though shallow, is relatively turbid and it is thought that stagnant conditions associated with low wind allow positively buoyant cyanobacteria to obtain favorable, near-surface light conditions. The combination of remote sensing, *in situ* biogeochemical studies, and local meteorological data allowed us to investigate these linkages over large spatial scales. Measurements of benthic and pelagic oxygen metabolism along with spatial patterns of MERIS-derived chl-a showed that 60–95% of the area of the

lagoon was vulnerable to transient hypoxia when blooms coincide with calm conditions (Zilius et al., 2014).

HYDRODYNAMIC FACTORS AS DRIVERS OF BLOOMS

Freshwater inputs to the lagoon are dominated by the Nemunas River, which has an annual average discharge of 21.8 km³, and accounts for 96% of total inputs (Jakimavičius and Kriaučiūnienė, 2013). The Nemunas River discharges into the central part of lagoon, dividing the system in a northern and a southern region that differ in water renewal time (Umgiesser et al., 2016). The northern part of the lagoon is characterized by strong riverine influence and short renewal time (< 80 days), which result in limited accumulation of suspended matter (Ferrarin et al., 2008; Remeikaitė-Nikienė et al., 2016). The southern part of the lagoon has a longer water residence time (> 190 days) with minimal fluvial influence. The latest efforts to analyze the water exchange within the Curonian Lagoon (Umgiesser et al., 2016) revealed different seasonal patterns of residence time, primarily driven by changes in hydrographic forcing by the Nemunas River. During elevated spring discharge, the entire lagoon is strongly flushed by Nemunas River. During summer, river discharge decreases, resulting in increased water residence time, particularly in the southern lagoon. Summer stagnation has implications for water temperature, stratification, nutrient availability and stoichiometry, and phytoplankton abundance and composition. In summer, wind forcing appears to be the most important factor influencing water column mixing and



exchange between the southern and the northern part of the lagoon (Umgiesser et al., 2016).

NUTRIENT LOADS AND THEIR ECOLOGICAL STOICHIOMETRY

A study coupling the Curonian Lagoon with its watershed was started in 2012 to better characterize the timing of nutrient inputs and their stoichiometry. From 2012, on at least a monthly basis and more frequent (weekly) during high discharge periods, discharge and water chemistry (including all dissolved and particulate forms of N, Si, and P) were monitored near the inflow of the Nemunas River to the lagoon (Vybernaite-Lubiene et al., 2017; Figure 1). Discharge and nutrient concentrations displayed strong seasonality. Nitrate and reactive Si concentrations decreased by two orders of magnitude from spring to summer (e.g., NO_3^- from > 300 μM to < 1, SiO_2 from > 200 to < 1 $\mu M)$ while reactive P concentrations showed comparatively smaller changes (from 0.2 to 4 μM). These seasonal patterns resulted in reduced DIN:DSi and DIN:DIP ratios, which shift the lagoon from an excess of N and Si in colder months, to P excess (DIN:DIP < 16) in warmer months (Figure 3). These findings support the hypothesis that cyanobacterial blooms are favored during summer by the limited



deviations (n = 6) are reported.

N and Si supply via riverine inputs (Pilkaitytė and Razinkovas, 2007).

Our analyses of nutrient loads showed that recent (2012–2016) N export from the Nemunas River basin is similar to historical data (1986–2002), whereas P loads have declined by

nearly 60 % as a result of sewage treatment plant improvements (Vybernaite-Lubiene et al., 2018). Despite reductions in P loads, the lagoon remains imbalanced with an excess of P relative to N, thereby favoring the growth of N-fixing cyanobacteria. Further P reductions are needed to promote- limitation or co- limitation and thereby diminish the dominance of cyanobacteria. It is also important to stress that despite strong N limitation in the lagoon, hyperblooms of cyanobacteria do not occur every summer due to the influence of other factors that regulate blooms.

SEDIMENTS AND BENTHIC PROCESSES AS DRIVERS OF BLOOMS

The distribution of sediment types in the Curonian Lagoon is determined by hydrodynamic factors and by contributions from autochthonous and allochthonous materials (Pustelnikovas, 1994; Ferrarin et al., 2008). Curonian Lagoon sediments include a broad spectrum of deposits, from sand-dominated in the northern (riverine-influenced) sector to silt-dominated in the southern (more lentic) area (Trimonis et al., 2003). Declines in external (riverine) loads during the transition from spring to summer enhances the importance of internal recycling from the benthic compartment as a nutrient source for pelagic primary production. Studies of sedimentary processes revealed a shift in dominant microbial processes and benthic fluxes from spring to summer (Zilius et al., 2012, 2014; Petkuviene et al., 2016). For example, net N₂ production suggests the dominance of denitrification over N-fixation during spring; however N2 fluxes are reversed during summer, suggesting net N import to the benthic compartment (Zilius et al., 2018).

Since 2009, oxygen penetration depth, total and diffusive sedimentary oxygen demand, pore water chemical environments, sedimentary pools and benthic fluxes were measured or calculated at sites representative of dominant sedimentary environments including littoral, pelagic transitional and confined zones (Zilius et al., 2012; Figure 1). In this turbid system, benthic photosynthesis was measurable only in shallow littoral illuminated sediments (~1 m depth) representing a minor fraction (5%) of the total lagoon surface (Benelli et al., 2018). Here, benthic algae oxidize the upper sediment layer and efficiently retain nutrients, thereby impeding regeneration to the water column (Zilius et al., 2012; Benelli et al., 2018). Deeper sites were always heterotrophic and their seasonal oxygen metabolism and nutrient regeneration was driven by water temperature and phytoplankton blooms; recently settled fresh phytoplankton resulted in significantly higher oxygen uptake, limited oxygen penetration in sediments (< 1 mm), and high rates of anaerobic to aerobic metabolism (Zilius et al., 2012, 2016).

The mechanisms underlying P release from sediments were analyzed in detail, as they contribute to lower inorganic DIN:DIP ratio in the water column and favor cyanobacteria (Zilius et al., 2014, 2015, 2016; Petkuviene et al., 2016). The distribution of sedimentary pools of P, Fe, Mn and S in the Curonian Lagoon was related to riverine influence; sandy sediments adjacent the Nemunas delta were oxidized and have a large geochemical buffer capacity against the effects of anoxia, with limited accumulation of free sulfide (Petkuviene et al., 2016). Muddy areas along the western and southern portion of the lagoon had chemically reduced sediments where the reduction of iron may result in large P release (Petkuviene et al., 2016). In manipulative experimental studies, simulated deposition of phytoplankton material, primarily composed of cyanobacteria, resulted in an increase of dissimilative nitrate reduction over denitrification and large methane production, but with limited reactive P release (Zilius et al., 2016). Experimental manipulations of intact cores, targeting short-term effects of anoxia, revealed that pools of detritial Ca bound P (> 70 % of inorganic P) and oxidized Fe and Mn, prevent or buffer redox-dependent reactive P release from sediments (Zilius et al., 2015). However, seasonal measurements of reactive P fluxes at sandy and muddy areas revealed large summer P release at muddy sites coinciding with the occurrence of cyanobacterial blooms, and the onset of hypoxia and anoxia in the water column (Zilius et al., 2014; Petkuviene et al., 2016). These events occurred under specific conditions during prolonged stable weather, with no wind and high water temperature. Benthic P release occurred when the oxidized pools of metals within sediments were exhausted and contributed to the imbalanced stoichiometry by further lowering the DIN:DIP ratio. P regeneration from sediment, despite occurring over short period, had a significant effect on the lagoon P budget, resulting in a large export of P (Petkuviene et al., 2016).

TOP-DOWN DRIVERS OF BLOOMS

Zooplankton

The shift from diatom- to cyanobacteria- dominated phytoplankton communities was accompanied by a decline in relative zooplankton grazing. During the spring diatom bloom, maximum consumption by zooplankton corresponded to 34% of NPP (324 μ gCL⁻¹d⁻¹), whereas during the summer cyanobacteria bloom grazing decreased to 8 % of NNP $(470 \ \mu gCL^{-1}d^{-1})$ (Figure 4). A similar pattern was observed in the southern part of the lagoon where zooplankton grazing declined from 60% of phytoplankton production during spring to 4 % in summer (Semenova and Aleksandrov, 2009). Despite reduced grazing rates, stable isotope studies show that cyanobacteria blooms support secondary production in a diverse group of benthic and pelagic consumers within the lagoon (Lesutienė et al., 2014). Our studies also show that cyanotoxins (microcystin) are found in tissues of fish and shellfish, indicating that cyanobacteria production supports higher trophic levels in this system (Bukaveckas et al., 2017).

Grazing by zooplankton may be an important driver of cyanobacteria bloom development. During winter, ciliate growth is limited by low biomass of phytoplankton. In the early spring, when small-sized phytoplankton are dominant, the ciliate assemblage was dominated by small naked oligotrichs and prostomatids. After the late spring diatom bloom, the ciliate assemblage shifted to medium sized tintinnids, which feed on the same nano-fraction of phytoplankton or/and heterotrophic flagellates as ciliates. The summer/autumn phase was characterized by increased taxonomical and functional



diversity of ciliates indicating exploitation of a wide size range of food. Small sized naked oligotrichs (*Strobilidium* spp.) and peritrichs (*Vorticella* spp.) (mainly bacterivorous ciliates) dominated in summer, indicating a shift from algal food to bacteria (Grinienė, 2013; Grinienė et al., 2016). The shift from large *Daphnia* to small-bodied *Chydorus sphaericus* coincides with the dominance of cyanobacteria (Gasiūnaitė and Razinkovas, 2004; **Figure 5**). *Chydorus* graze on smaller algae and therefore give an advantage to large cyanobacteria (Gasiūnaitė and Olenina, 1997). In addition, the presence of large filamentous colonies and toxic strains may foster the dominance of bloom forming cyanobacteria (Pilkaitytė and Razinkovas, 2007).

Macrofauna

Excluding the littoral zone, sediments of the Curonian Lagoon host few macrofauna species due to high organic content and poor oxygen conditions (Zettler and Daunys, 2007). Among them, oligochaetes, chironomid larvae and freshwater mussels, including native unionids and invasive dreissenids, are dominant groups (Daunys, 2001). Chironomid larvae and mussel aggregations may, due to their high densities, influence phytoplankton composition and abundance (Dame et al., 1980; Officer et al., 1982; Gili and Coma, 1998). We discuss here if and under which circumstances macrofauna may favor the onset of cyanobacterial blooms in the Curonian Lagoon.

In the lagoon, periods of short water residence time may impede efficient removal of particulate matter by suspension feeding. In spring (average residence time 7 days) only 10 % of particulate matter was removed by zebra mussels, while in summer (average residence time up to 15 days), the proportion of particulate matter removed increased to 30% (Daunys et al., 2006). Chironomid larvae and mussels may exert a top-down control of pelagic primary production but



they may simultaneously excrete large amount of nutrients. Their activities also enhance the organic matter content of sediment via biodeposition, stimulating microbial activity and re-mineralization (Caraco et al., 1997; Stief, 2013; Ruginis et al., 2014; Benelli et al., 2017). It is unclear whether the net effect of phytoplankton removal via grazing is offset by nutrient regeneration via excretion and whether these processes have a specific benefit to cyanobacteria. Dreissena polymorpha was intensively studied due to its top-down control on phytoplankton and the possible management of its biomass to reverse eutrophication. However, such top-down control on pelagic primary production resulted to be site-specific and context-dependent (e.g., in shallow, well-mixed environments with low nutrient background more than in deep, stratified ecosystems with high nutrient inputs) (Conroy et al., 2005; Caraco et al., 2006). Furthermore, dreissenids excrete large amounts of reactive P and different authors have suggested that these mussels may change nutrient stoichiometry, via P mobilization and by enhancing N removal via denitrification (Zhang et al., 2008; Ruginis et al., 2014). The inability of zebra mussels to graze on larger forms of cyanobacteria may provide a competitive advantage over other algae, which, in combination with increased rates of reactive P re-cycling, enhances the potential for cyanobacteria blooms. These aspects need further study, but suggest that the presence of dreissenids mussels on the Curonian Lagoon may exacerbate the effects of nutrient loading, and favor increased dominance by cyanobacteria.

Birds

The Curonian Lagoon hosts a large bird community, including tufted ducks and common pochards with 24,500–54,700 and 1,800–41,000 individuals, respectively (Stanevičius et al., 2009),

goosanders (Žydelis, 2001), cormorants, with more than 10,000 breading birds (Švažas et al., 2011; Dagys and Zarankaitë, 2013), mallards, geese (3,000–6,500 ind/day) and little and blackheaded gulls (1,000–1,500 ind./day). High densities of water birds are vectors of seeds, invertebrates, bacteria and phytoplankton (Tobiessen and Wheat, 2000), and also contribute to nutrient loads (Manny et al., 1994; Hahn et al., 2007; Green and Elmberg, 2014; Han et al., 2017).

During nesting, breeding and roosting periods, water birds enrich the water with guano (Klimaszyk et al., 2015). In enclosed aquatic ecosystems bird feces may contribute 50–69%, 27– 40%, and 70–75% of total C, N, and P loads, respectively (Manny et al., 1994; Post et al., 1998; Boros et al., 2008; Gwiazda et al., 2014). Bird feces have low N:P, implying that water bird excretions may strengthen N limitation and promote cyanobacteria blooms (Rönicke et al., 2008; Han et al., 2017). Birds also have indirect effects on nutrient cycling by removing macrophytes, invertebrates and fish. Herbivorous birds, by removing plants, remove those elements that trap nutrients in the benthic compartment, provide shelter for zooplankton and allow sediment oxidation via radial oxygen loss. While grazing on macrophytes, birds resuspend sediments and mobilize pore water nutrients (Klimaszyk et al., 2015; Klimaszyk and Rzymski, 2016). Furthermore, a large fraction of macrophyte-associated P is released to the water column in the reactive form and is readily available to phytoplankton. The mechanisms that regulate P mobility in sediments are redox-dependent such that the removal of roots and macrofauna, together with particle resuspension, has the potential to mobilize sediment sources of P. In the Curonian Lagoon herbivorous birds represent the second largest water bird group, peaking in spring and distributed throughout the Nemunas River deltaic area and the littoral zone.

Benthivorous birds, feeding on macrofauna, produce an effect on the benthic system similar to that of fishes, removing animals that may keep the sediment oxidized and resuspending sediments and nutrients (Werner et al., 2005; Sánchez et al., 2006; Rodríguez-Pérez et al., 2007; Matuszak et al., 2014). Piscivorous birds convert fish-associated P into reactive P (P \bar{u} tys and Zarankaitė, 2010). Large colonies of cormorants have their peak activity during summer. The large bird community in the Curonian Lagoon may therefore affect by various direct and indirect mechanisms the cycling of nutrients and that of P in particular.



Fish

Benthivorous fish, including carp, roach, bream and perch, represent the dominant fish component in the Curonian Lagoon (Cline et al., 1994; Persson and Svensson, 2006; Lithuanian Environmental Protection Agency (EPA), 2008; Adámek and Maršálek, 2013). Fish may produce both top-down (e.g., removal of grazers and competitors) and bottom-up effects (nutrient mobilization) that favor eutrophic conditions and cyanobacterial blooms (Shormann and Cotner, 1997; Roozen et al., 2007). The diet of benthivorous fish in the Curonian Lagoon includes mussels, chironomidae larvae, detritus, zooplankton and plants (Bubinas and Ložys, 2000). Benthivorous fish may impact the water quality, leading to nutrient accumulation and algal growth, by suspending the sediments and by feeding on filterfeeding zooplankton, burrowing macrofauna and macrophytes (Zambrano and Hinojosa, 1999; Williams et al., 2002; Parkos et al., 2003). Sediment resuspension by the benthic fish community increases water turbidity, limits light penetration and rooted macrophytes and favors P release and cyanobacteria growth (Hellström, 1991; Breukelaar et al., 1994). By removing invertebrates from the sediments, benthivorous fishes mobilize nutrients from the pore water (Tarvainen et al., 2002; Phan-Van et al., 2008). Resuspension itself may oxidize sediments, but this is a short-term and local effect, while reductions in invertebrate abundance impacts N removal via denitrification and P sequestration. Moreover, fish predation reduces zooplankton populations, resulting in low grazing on phytoplankton (Jeppesen et al., 1999). Fish excretions are very soluble and rich in N and P which stimulate periphyton growth and negatively affect macrophytes (Tarvainen et al., 2002; Williams et al., 2002). Excreted nutrients are dispersed horizontally and vertically and from littoral to pelagic areas (Schindler et al., 1996; Persson and Svensson, 2006).

SYNTHESIS

Cyanobacterial blooms in the Curonian Lagoon arise from multiple interacting factors, which include external forcing (riverine discharge and wind conditions) and internal processes (consumer-mediated nutrient cycling and sediment-water nutrient exchange). We summarize the information discussed in this review through a graphical representation of the multiple mechanisms that drive cyanobacterial blooms in the Curonian Lagoon (**Figure 6**).

During spring, the lagoon is diatom-dominated due to a combination of low water temperatures, high river discharge and availability of inorganic N and Si, in excess to P. The system alternates phases with clear and turbid water depending on the intensity of the spring diatom bloom and the occurrence of wind-associated sediment resuspension events. During spring, light penetration may attain 1–2 m and grazers exert appreciable control of algal biomass accrual. The water column is generally well mixed and normoxic; under these circumstances the upper sediment layer is oxidized and acts as a nutrient sink.

The spring-summer transition is marked by a decline in discharge of the Nemunas River, which is accompanied by

the depletion of N and Si within the lagoon. Reductions in external loadings, together with processes occurring within the lagoon (high spring denitrification rates and Si sequestration via uptake and accumulation in sediment), result in the onset of N and Si limitation. Cyanobacteria become dominant, resulting in a series of cascade effects that include increased algal-associated turbidity and water stratification. Positive feedbacks arise as large colonies of cyanobacteria limit the capacity of grazers to control biomass accrual and high respiration rates promote oxygen undersaturation. Climatic conditions, which are highly variable, play an important role, as low wind conditions may further push the system toward hyperbloom events, with extensive surface scums (Figure 7). Water column respiration, not sediment oxygen demand, promote oxygen depletion in the system, due to the large availability of labile organic matter from decaying algal cells. By this mechanism, hyperblooms promote their persistence as hypoxia results in sediment P release. Other factors that may contribute to these large periodic outbreaks include the presence of waterbirds. Large colonies of cormorants settle in spring along the Curonian Spit and have their most intense period of activity during summer. Cormorants, through the production of guano, make large amounts of P available in surface water. Besides cormorants, colonies of seagulls, swans and duck have large numbers and may contribute to make fish or macrophyte P pools readily available to cyanobacteria. Other biological agents supporting algal blooms include the invasive freshwater mussel (D. polymorpha), which excretes large amounts of reactive P relative to the native unionid mussels and may therefore contribute to the low DIN:DIP ratio in the lagoon.



FIGURE 7 | Cyanobacteria hyperbloom with scum formation in the Curonian Lagoon, August 2013.

MANAGEMENT POLICIES AND PERSPECTIVES

The Curonian Lagoon ecosystem provides a number of provisioning and cultural ecosystem services, most directly linked to the main economic activities in the lagoon area - recreation and fishery (Rashleigh et al., 2012; Razinkovas-Baziukas et al., 2012). The lagoon also provides ecosystem services of relevance to the Baltic Sea region such as denitrification and phosphorus burial. Management efforts to improve water quality in the Curonian Lagoon have targeted reductions of nitrogen loads by 15 % and phosphorus by 8%. Model simulations (Ertürk et al., 2016) revealed that reductions of nitrogen loads by 14 % and phosphorus loads by 6%, will bring about a 10% reduction in the abundance of cyanobacteria (Razinkovas et al., 2008). Further reductions in riverine nutrient loads (40% decrease in both N and P) produced only a 10% decrease in peak chl-a concentrations (Razinkovas et al., 2008). Further efforts to improve water quality may require withinsystem bioengineering solutions (biomanipulations, mussels, reed harvesting).

Climate change projections for the Curonian Lagoon (Jakimavičius et al., 2018) indicate an increase of average water temperature up to 1.7-2°C by the middle of this century, consistent with trends observed during the last three decades (Dailidienë et al., 2011). According to this modeling study, the increase in water temperature was mostly confined to the summer-early autumn period, which may therefore favor the development of cyanobacteria blooms. Biogeochemical cycles of the Curonian Lagoon will be affected by changes in the water balance of the lagoon. A decline in contributions from Nemunas River coupled with an increase in Baltic water intrusions (due to sea level rise) will alter the water balance during the winter-spring period. A shift in the timing of peak discharge from spring to winter, as observed in recent decades (Dailidienë et al., 2011), may diminish algal blooms if a larger proportion of the nitrogen load from the Nemunas River passes through the Curonian Lagoon during the period of low phytoplankton productivity. However, the predicted decrease of ice cover is expected to reduce winter hypoxia, which would result in reductions in denitrification. Despite the intensive studies of this system, we are not able to predict whether climate change will exacerbate or mitigate cyanobacteria blooms. However, it is apparent that further management actions are needed to reduce nutrient loads and restore ecosystem services. There is a need for additional studies, both at the watershed-scale and the lagoon scale, to facilitate science-based management decision. At the watershed scale, long-term monitoring is needed to better understand the effectiveness of improved agricultural practices and water treatment on N, Si, and P export from the Nemunas basin to the Curonian Lagoon. Watershed practices may differentially affect the three elements further modifying their ecological stoichiometry, with implication for algal blooms

(Yunev et al., 2007; Bresciani et al., 2014; Vybernaite-Lubiene et al., 2017).

CONCLUSION

This review analyzes the available information on the mechanisms driving cyanobacterial blooms in the Curonian Lagoon. Results from our analysis suggest that blooms are a consequence of multiple, interplaying factors, producing a cascade of processes and positive feedbacks. The hot moment for cyanobacteria blooms is the summer, due to combination of favorable nutrient stoichiometry (N and Si limitation), elevated water temperature, low wind speed, unbalanced internal recycling (P > N) and low grazing pressure. The hot spots of cyanobacteria are stagnant areas where limited water circulation and stratification provide these organisms a competitive advantage. These hot spots may serve as bloom initiation areas from which cyanobacteria are dispersed by prevailing winds. Ecological interactions among aquatic organisms, and how these respond to changes in climate and to species invasions remain understudied. The combination of satellite remote sensing, traditional monitoring of environmental parameters, detailed analysis of processes at the macro and microscale and the application of ecological network models, have proved to be useful tools for understanding the mechanisms underlying the development of cyanobacteria blooms. Our further efforts seek to improve our capacity to predict the occurrence and severity of algal blooms and guide prevention measures.

AUTHOR CONTRIBUTIONS

MBa collected and homogenized the material with the help of MZ, who realized the conceptual scheme of **Figure 6** and PB, who organized the different sections and improved the clarity of the paper. All authors contributed to the writing.

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