A 45-year hydrological and planktonic time series in the South Bight of the North Sea

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Abstract. This article presents a 45-year data series (from 1978 to 2023) acquired under the IGA (Impact des Grands Aménagements in French, Impacts of Major Developments in English) program conducted by IFREMER for EDF (Électricité de France, the French multinational electricity utility company). The IGA program was established to monitor the ecological and environmental quality of the coastal area surrounding the Gravelines Nuclear Power Plant (GNPP) located in the South Bight of the North Sea. The main objective of this program is to assess medium- and long-term environmental evolution by providing the means to identify possible changes in local marine habitats. Since 1978, the IGA program has measured key parameters, including temperature, salinity, nutrient concentrations, oxygen levels, chlorophyll-a concentrations, and the abundance of phytoplankton and zooplankton species. These measurements have been taken at different sampling stations around the GNPP, including the Canal d'amenée sampling station, for which hydrological and biological characteristics are considered representative of the broader coastal area of the South Bight of the North Sea. This data paper provides an overview of the main statistical characteristics of the time series (available at https://doi.org/10.17882/102656, Lefebvre et al., 2024), including long-term trends and shift analysis. Despite the importance and length of this dataset, one of the longest available for this region, its application in advancing knowledge of hydrological and biological processes has been surprisingly limited. The aim of this paper is to make this valuable dataset available to the scientific community, stakeholders, and society to help decipher the local and global influences of anthropogenic activities in a world increasingly affected by climate change. Since all the main statistics and patterns are still available thanks to our analysis, users should be able to use these data and combine them with other sources (in situ, satellite, and modeling) in order to dive into deeper analyses and investigate new key scientific challenges and more specific ones.

1 Introduction

Understanding the dynamics of marine ecosystems requires long-term time series that allow us to take into account different sources of natural variability (seasonal and interannual) and anthropogenic variability (under local or regional pressures and global warming). However, such long-term series are difficult to obtain, as it is costly to maintain and require measurement protocols that are stabilized throughout their duration to avoid acquisition biases. Long-term planktonic time series, particularly for phytoplankton and zooplankton, are especially relatively scarce and not widely available in the scientific literature. Phytoplankton form the basis of pelagic food chains and are responsible for approximately 50% of global primary production (Falkowski et al., 2003), playing a crucial role in capturing carbon dioxide (CO₂) and producing oxygen (O₂). Under certain favorable conditions of temperature, light, turbulence, and nutrient availability, the growth of various phytoplankton species can be enhanced. Among the species forming massive blooms in the eastern English Channel and South Bight of the North Sea (SBNS), some, like *Pseudo-Nitzschia, Dinophysis*, and *Alexandrium*, are known to produce phycotoxins that pose significant health risks to humans when introduced into the food chain, particularly through filter-feeding shellfish. Other species, such as *Phaeocystis globosa* (Lefebvre and Dezécache, 2020; Karasiewicz and Lefebvre, 2022; Karasiewicz et al., 2018; Lancelot and Rousseau, 1994), can become so abundant that they disrupt ecosystem functioning. Both physicochemical parameters (bottom-up control) and zooplankton populations (top-down control) influence the phenology of phytoplankton species (Banse, 1992; Feng et al., 2014).

Zooplankton form the main aquatic animal compartment in terms of biomass and diversity (Mauchline, 1998). Zooplankton are considered primary consumers, as they are major consumers of phytoplankton (Atkinson, 1996), which makes this organic matter available to predatory animals, particularly fish larvae and juveniles. In the eastern English Channel and SBNS, holoplankton (i.e., species that stay planktonic throughout their life cycle, in contrast to meroplankton) are dominated by copepod species such as the calanoids Temora longicornis, Acartia clausii, and Centropages hamatus and the harpacticoid Euterpina acutifrons (in order of annual phenology and dominance; Brylinski, 2009). Zooplankton communities are generally studied on an ad hoc basis, and long-term zooplankton time series are even rarer in the scientific literature than phytoplankton time series, especially in the eastern English Channel and the French part of the SBNS.

Ecological monitoring of nuclear power plant (NPP) discharges into the sea focuses on studying the medium- to long-term temporal evolution of various marine domains, including the pelagic, benthic, and fishery domains, along with their associated compartments (hydrology, physicochemistry, chemistry, phytoplankton, zooplankton, benthos, and microbiology). This monitoring is conducted on a localized spatial scale, focusing on areas surrounding the power plants and within their zones of influence. The aim is to detect any changes from monitoring specific parameters that are characteristic of each marine compartment. Ecological monitoring of such infrastructure in Gravelines, located in the French coastal part of the Dover Straight, was initiated in 1978, providing the longest-running ecological time series in this part of the English Channel-North Sea continuum. The monitoring strategy and sampling methods (including the sampling locations and frequencies and the nature of the analyses carried out) evolved between 1978 and 1986 as the six production units of the Gravelines power plant came on stream. One of the monitoring stations (named Canal d'amenée) of this network was specifically set up to track the characteristics of seawater entering the cooling turbines and is therefore not influenced by the NPP activity. This monitoring station, crucial for smooth operation of NPPs, has benefited from an improvement of the monitoring protocol by increasing the frequency of the measurements. Particular attention has been paid to plankton monitoring, where certain species (gelatinous zooplankton or harmful algal bloom species (HABs) such as Phaeocystis globosa) are known to cause seasonal disruptions to NPP operations (Masilamoni et al., 2000; Wang et al., 2022, 2023). Initially established to assess the direct and indirect effects of large-scale coastal developments, this monitoring effort now also offers scientists a valuable opportunity to study local plankton dynamics on a multidecadal scale.

2 **Objectives**

The aim of this paper is to present the IGA (Impact des Grands Aménagements in French, Impacts of Major Developments in English) physicochemical, phytoplankton, and zooplankton dataset at the Canal d'amenée sampling point. This includes an overview of the sampling strategy, data collection process (with the associated quality-assurance and quality-control steps), data investigation, and storage. The characteristics of the different datasets and a general interpretation of their variability will be detailed. Based on the limited existing applications and valorizations of the IGA dataset, we will demonstrate its relevance not only for furthering understanding of marine phytoplankton ecology, but also public policy needs, such as assessment of environmental or ecological status as requested by EU directives and regional sea conventions. Additionally, we introduce some numerical tools based on an R package available for the scientific community and developed specifically to rapidly process such data and therefore valorize the findings.

3 Materials and methods

3.1 Sampling point location

The GNPP site is located in the South Bight of the North Sea, near the Dover Strait. This location, coupled with the shallow waters (less than 30 m), results in strong hydrodynamic conditions. The hydrodynamic regime of the southern North Sea is influenced by the semidiurnal tidal circulation and wind action. The range is macrotidal, varying from 3.5 m during neap tides to 5.6 m during spring tides. Tidal currents along the Dunkirk coast run parallel to the coast and are asymmetrical: the flood tide, flowing northeast, is faster and more intense than the ebb tide, which flows southwest. This flood current dominates the entire southern North Sea coastline (Fig. 1), with recorded maximum speeds of approximately 1.5 m s^{-1} . Off Dunkirk, "the flood is present between 3 h before and 3 h after high tide" (SHOM, 1988). The only river flowing into the North Sea along the French coast is the Aa, a river 89 km long with a watershed of 1215 km² and a discharge rate of $10 \text{ m}^{-3} \text{ s}^{-1}$. Industrial activities represent the main local pressures on this environment.

The most comprehensive ecological time series are available from the Canal d'amenée sampling station (x: 2.15003, y: 51.0222), located within the Gravelines harbor at the entrance of the NPP cooling system (Fig. 1). This position allows the station to avoid the direct impacts of the NPP operations, although it remains subject to other anthropogenic in-



Figure 1. Location of the Canal d'amenée sampling station (black circle) in the Gravelines harbor, which is open on the South Bight of the North Sea.

fluences such as global warming and eutrophication. This is therefore a relevant position for coastal monitoring, completing the monitoring network of the French North Sea coastal ecosystem alongside the SRN Dunkerque 1 station (Lefebvre and Devreker, 2023).

3.2 Hydrological and biological parameter measurements

Water samples were initially collected at a monthly frequency and later at a weekly frequency at the end of the 1980s at the surface of the Canal d'amenée sampling station. Water samples were analyzed for different parameter measurements, though not all parameters were measured at every sampling. The evolution of the monitoring practices and the specific temporal distribution of the samples are detailed by the parameters in the Supplement (Table S1 and Figs. S1– S3).

3.2.1 Temperature, salinity, turbidity, and oxygen concentration

These four variables are measured using a multiparameter probe. Salinity is determined using a conductivity sensor coupled with a temperature sensor (°C) and is expressed as a conductivity ratio (PSU for practical salinity unit). Unlike temperature measurements, which conform with the global frequency strategy (i.e., monthly sampling until 1985 and weekly since), salinity was recorded only twice a year between 1990 and 2006 and has been monitored weekly since 2013.

Since 2019, turbidity has been assessed by nephelometry and expressed in nephelometric turbidity units (NTUs) (USEPA, 1980), but it was measured with a FNU (formazine nephelometric unit, ISO 7027) sensor between 2016 and 2018. Turbidity is quantified by measuring the amount of light scattered at 90° compared to the incident light. Dissolved oxygen concentration has been measured using a luminescence oxygen sensor since 2016 (mg L⁻¹) (NF EN ISO 25814). However, prior to 1987, it was measured using the Winkler method (mL L⁻¹). The complete methods are described in Aminot and Kérouel (2004). Oxygen concentrations have only been measured in the 1970s, in the 1980s, and since 2016.

Except for turbidity, these variables are recognized as essential ocean variables (EOVs) for physics and biochemistry by the Global Ocean Observing System (GOOS) (Muller-Karger et al., 2018). 1176

3.2.2 Nutrients

For nutrient analysis, a 65 mL subsample (for phosphate and dissolved inorganic nitrogen DIN = ammonium + nitrite + nitrate) or a 125 mL subsample (for silicate) of water filtered through a 48 μ m mesh is frozen at -25 °C. Nutrient concentrations are subsequently measured in the laboratory using a spectrophotometer with an optical-density (OD)-concentration relationship determined from a calibration curve performed for each series of measurements. The complete methods are detailed in Aminot and Kérouel (2007).

Ammonium (NH₄⁺) is quantified using the indophenol blue molecular absorption spectrophotometric method (NF T90-015-2; μ mol L⁻¹). Nitrite (NO₂⁻) and nitrate (NO₃⁻) are measured using the N-naphthyl-ethylenediamine molecular absorption spectrophotometric method (NF ISO 13395; μ mol L⁻¹). Phosphate (PO₄²⁻) is determined by the phosphomolybdic blue molecular absorption spectrophotometric method (NF ISO 6878; μ mol L⁻¹). Silicate (SiOH) in water is measured using the silico-molybdic blue molecular absorption spectrophotometric method (NF T90-007; μ mol L⁻¹).

Nutrient monitoring was suspended in the mid-1990s and 2000s for nitrates and nitrites. Phosphate measurements were more intermittent, conducted only from 1978 to 1986 and then resumed from 2016 to the present at a monthly frequency. The Global Ocean Observing System also classifies nutrients as essential ocean variables for biochemistry.

3.2.3 Chlorophyll a

To measure chlorophyll pigments, an indicator of phytoplankton biomass, 1 L of water is kept cool and protected from light. To determine the specific composition and abundance of phytoplankton, 500 mL of water is fixed with Lugol's solution (2.5 mLL^{-1} of seawater). Until 2018, chlorophyll-*a* concentrations were determined using the trichromatic method (SCOR-UNESCO, 1966). From 31 December 2018, the measurement method used has been the monochromatic method (Lorenzen, 1967). Regardless of the method, 1 L water samples are filtered through Whatman GF/C 47 mm glass-fiber filters. The chlorophyll pigments concentrated in these filters are then extracted with 90 % acetone. After centrifugation for 20 min at 6000 rpm, the absorbance of the supernatant is measured spectrophotometrically. The limit of quantification is 0.10 µg L⁻¹.

Chlorophyll-*a* concentration (a proxy of phytoplankton biomass) is measured according to the global frequency strategy and is recognized as a Biology and Ecosystems EOV by the GOOS.

3.3 Plankton

3.3.1 Phytoplankton

A 10 mL volume of the Lugol-fixed water samples is decanted in a sedimentation tank for at least 12h, following the method of Utermöhl (1958). Cell counts are then performed using inverted microscopy within a month of sample collection to minimize significant changes in phytoplankton size and abundance. Except for Phaeocystis globosa enumeration, over 200 phytoplankton cells per sample are counted using a 20X Plan Ph1 0.5NA objective, yielding an error margin of 10%. Therefore, microphytoplankton (cell sizes between 20 and 200 µm) were mainly identified. For P. globosa, only the total number of cells is computed. A minimum of 50 solitary cells are enumerated from several randomly selected fields (10 to 30) using a 40X Plan Ph2 0.75NA objective. The abundance of cells in a colony is determined using a relationship between colony biovolume and cell number, as defined by Rousseau et al. (1990). No counting data for P. globosa are available between 1982 and 1990 for unknown reasons.

Phytoplankton identification is standardized using the WoRMS (2024) database and reaches the species level in many cases (Table 1). However, when identification is challenging or uncertain, a lower taxonomic level is kept. Some species are also grouped into "artificial taxa", "complex" species, or common higher taxonomic ranks (mainly genus or family) if they are subject to strong identification confusion (this is the case for Pseudo-nitzschia or Chaetoceros, for example). In such cases, "[]" denotes a complex of species, while "+" denotes a complex of species and a common genus. These groupings are considered taxonomic units in order to maintain consistency in the database despite changes in taxonomic names. Freshwater phytoplankton and protozoa, though regularly found in low abundances in samples, are not considered in figures and statistical analyses that only focus on marine species. Phytoplankton species diversity calculations and the Shannon-Wiener index (Eq. 1) are based on all the taxonomic levels:

$$H' = -\sum_{i=1}^{S} p_i \ln p_i,$$
 (1)

with H' the index value and p_i the proportion of the entire community made up of species *i*.

During the period from 1978 to 2023 (excluding 1987, which is a missing year), a total of 1811 phytoplankton samples were collected, representing 237 taxa across various taxonomic ranks, which are mostly species and genera, providing good overall taxonomic accuracy (Table 1). Of these taxa, the Prymnesiophyceae species *Phaeocystis globosa* is the most abundant, with blooms reaching "more than millions" of cells per liter each year. The six other prevalent taxa are Bacillariophyceae species (i.e., *Rhizosolenia delicatula, Chaetoceros*, and *Pseudo-nitzschia [calliantha + delicatissima + pseudodelicatissima + subcurvata]*; *Rhizosolenia* sp.; *Skeletonema costatum*; and *Leptocylindrus [danicus + cur-*

Taxonomic rank	Number	Occurrence		
Kingdom	1	2		
Infraphylum	1	209		
Phylum	3	934		
Form	2	4		
Class	6	1243		
Order	5	928		
Family	7	99		
Complex	6	1273		
Genus	107	14 740		
Species	112	17 972		

Table 1. Number and occurrence of the different taxonomic ranks (highest level of taxonomic identification) encountered in phytoplankton identification.

vatus + *mediterraneus* + *aporus* + *convexus* + *hargravesii* + *adriaticus*]). The seventh most common taxa are Cryptophyceae, while the first Dinophyceae genus is *Gymnodinium*, which is ranked in 37th position. However, the relative dominance of these taxa can vary with the seasons and years.

Phytoplankton sampling frequencies conformed to the global frequency strategy, and phytoplankton diversity is recognized as a Biology and Ecosystems EOV by the GOOS.

3.3.2 Zooplankton

Zooplankton sampling at the Canal d'amenée station was conducted monthly using a WP2 net equipped with a flow meter for water volume calculations of each sample. By using a WP2 net, mesozooplankton (0.2-2 cm) and larger planktonic organisms were mainly sampled. The sampling duration was 10 min for gelatinous zooplankton and 3 min for mesozooplankton. Over time, the type of plankton net used has changed. From 1991 to 2007, the standard WP2 net, used since the start of monitoring in 1978, was replaced with a smaller net $(0.09 \text{ m}^2 \text{ opening}, 110 \text{ cm high, tapered, and with})$ a 200 µm mesh size) for sampling in the intake channels. The smaller opening and conical shape of this net made it more sensitive to the effects of clogging and backflow, particularly in spring and summer, when algae (Phaeocystis globosa and Coscinodiscus sp.) or Noctiluca proliferate. The WP2 net, with its wider opening and cylindrical-conical shape, prevents backflow into the net and is more effective for capturing zooplankton from a single point. From 2008 onward, the standard WP2 net was reintroduced for sampling. A 2018 analysis of potential biases related to changes in sampling strategies (frequency, net type, and level of identification) indicated that the replacement of the smaller net with a WP2 net did not significantly impact the estimates of the total zooplankton abundance.

Samples are preserved in formalin solution (0.9% final concentration). They are filtered through a 200 µm mesh at the laboratory, and subsamples are obtained using a Motoda

Table 2. Number and occurrence of the different taxonomic ranks (highest level of taxonomic identification) encountered in zooplankton identification.

Taxonomic rank	Number	Occurrence		
Kingdom	1	87		
Subclass	2	503		
Infraorder	4	393		
Phylum	8	969		
Order	12	1136		
Class	12	1288		
Family	24	760		
Genus	53	1226		
Species	108	6091		

box and identified under a binocular microscope in a Dolfus tank. Identification stops when 100 individuals are counted. The number of individuals counted is then extrapolated to the total sample volume (ind m⁻³). Zooplankton identification is standardized using the WoRMS (2024) database, achieving species-level identification in many cases (Table 2). However, as for phytoplankton, when identification is challenging or uncertain, a lower taxonomic level is retained. To prevent misinterpretation of scientific names, the AlphaID, which comes from the WoRMS reference website, is also provided. Species richness calculations with the Shannon– Wiener index (Eq. 1) are based on all taxonomic levels (not only species).

During the 1978–2023 period, 585 zooplankton samples were collected, representing 224 taxa across different taxonomic ranks that are mostly species and genus and providing a high taxonomic accuracy. The most abundant species identified is the calanoid copepod *Temora longicornis*. Copepods are globally the most abundant zooplankton taxa (*Acartia clausi, Euterpina acutifrons, Centropages hamatus, Pseudocalanus elongatus*, and *Paracalanus parvus*). Additionally, the appendicular *Oikopleura dioica* is frequently observed. However, this relative dominance varies between seasons and years. Zooplankton diversity is identified as a Biology and Ecosystems EOV by the GOOS.

3.4 Other parameters

The IGA survey monitor adds parameters, but these were only measured sporadically over a few years, depending on the monitoring strategy. These parameters, including pheopigments, zooplankton biomass, pH, suspended organic matter, nitrogen, and carbon organic concentrations in zooplankton, are scattered across the survey period and have therefore not been included in this article.

Table 3. Statistical summary (minimum, first and third quantiles, mean, median, maximum, and length of the data series) of the main zooplankton and phytoplankton taxonomic groups and physicochemical variables collected within the IGA monitoring program (1978–2023) at the Canal d'amenée station. The "Trend" column indicates whether there is an increasing (orange arrow) monotonic trend, a decreasing (green arrow) monotonic trend, or no significant trend ("na"). The "Trend %unit yr⁻¹" column provides a quantification of significant trends (as a percentage change per year using the Theil–Sen slope method).

$\begin{tabular}{ c c c c c c c c c c c c c c c c c c c$	Taxonomic group	Rank	Trend	Trend	Min.	1st quantile	Median	Mean	3rd quantile	Max.	Ν
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Paracalanus Pseudocalanus Genus Genus Genus Genus Guild ▲ 0.06 0.10 9.21 25.10 85.70 76.66 1447.22 436 Meroplankton total Guild -0.03 0.10 7.44 25.16 96.15 84.24 2954.50 397 Meroplankton total Guild -0.01 0.01 34.07 191.18 477.08 535.06 11818.61 578 Phytoplankton (cell L ⁻¹) - Bacillariophyceae Class na 500 33 300 110 850 274.944 289.475 21 669.833 1807 Pseudo-nitzschia Genus na 100 1900 8700 68.557 36.726 5340.000 1387 Pseudo-nitzschia Genus na 100 2200 5262 10539 11 550 919.000 1193 Rhizosolenia Genus na 100 2200 5500 30135 19.294 2249100 897 Thlassiosira + Porosira Genus na 30 800	Euterpina	Genus	na		0.10	13.30	59.70	315.30	301.30	8963.60	437
Pseudocalanus Meroplankton total Genus Guild ✓ −0.03 −0.01 0.10 0.01 7.44 34.07 25.16 191.18 96.15 477.08 84.24 535.06 2954.50 11818.61 397 Phytoplankton total Guild ✓ −0.01 0.01 34.07 191.18 477.08 535.06 11818.61 578 Phytoplankton (cell L ⁻¹) 500 33 300 110 850 274.944 289.475 21 669.833 1807 Chaetoceros Genus na 10 1900 8700 68.557 36 726 5 340.000 1387 Pseudo-nitzschia Genus na 100 2200 5262 10539 11 550 919.000 1693 Rhizosolenia Genus na 100 2200 5500 30 135 19 294 22 49 100 897 Thlassiosira + Porosira Genus na 30 800 2477 14 597 7893 1500.000 1595 Leptocylindrus Genus A 0.07	Paracalanus	Genus		0.06	0.10	9.21	25.10	85.70	76.66	1447.22	436
Meroplankton total Guild ▼ -0.01 0.01 34.07 191.18 477.08 535.06 11818.61 578 Phytoplankton (cell L ⁻¹) Bacillariophyceae Class na 500 33 300 110850 274 944 289 475 21 669 833 1807 Chaetoceros Genus na 10 1900 8700 68 557 36 726 5 340 000 1387 Pseudo-nitzschia Genus A 0.04 66 1300 6139 47 403 28 064 275 3000 1300 Paralia Genus na 100 2200 5262 10 539 11 50 919 000 1300 Skeletonema Genus na 100 2200 5500 30 135 19 294 2 249 100 897 Thisssissira + Porosira Genus na 100 2600 9360 68 055 36 130 11 886 000 879 Asterionella + Asterionellopsis Genus 0.03 100 1500 <td< td=""><td>Pseudocalanus</td><td>Genus</td><td>•</td><td>-0.03</td><td>0.10</td><td>7.44</td><td>25.16</td><td>96.15</td><td>84.24</td><td>2954.50</td><td>397</td></td<>	Pseudocalanus	Genus	•	-0.03	0.10	7.44	25.16	96.15	84.24	2954.50	397
Phytoplankton (cell L^-1)BacillariophyceaeClassna50033 300110 850274 944289 47521 669 8331807ChaetocerosGenusna101900870068 55736 7265 340 0001387Pseudo-nitzschiaGenus \blacktriangle 0.04661300613947 40328 0642 753 0001300ParaliaGenusna1002200526210 53911 550919 0001193RhizosoleniaGenus \checkmark -0.021001600800067 81144 1009 970 0001619SkeletomemaGenusna1002200550030 13519 2942 249 100897Thlassiosira + PorosiraGenusna30800247714 59778931 500 0001595Leptocylindrus + TenuicylindrusGenus \triangle 0.071002600936068 05536 13011 886 000870Eucampia + ClimacodiumGenus \frown -0.0280800245016 88384662 530 000920CryptophyceaeClass \triangle 0.191002100701621 62920 171349 600824DictyochophyceaeClass \triangle 0.030.0220057788777016299DinophyceaeClass \triangle 0.030.22005788777016299DinophyceaeClass \triangle 0.03<	Meroplankton total	Guild	•	-0.01	0.01	34.07	191.18	477.08	535.06	11818.61	578
Bacillariophyceae Class na 500 33 300 110 850 274 944 289 475 21 669 833 1807 Chaetoceros Genus na 10 1900 8700 68 557 36 726 5 340 000 1387 Pseudo-nitzschia Genus A 0.04 66 1300 6139 47 403 28 064 2 753 000 1300 Paralia Genus na 100 2200 5262 10 539 11 550 919 000 1193 Rhizosolenia Genus na 100 2200 5503 30 135 19 294 2 249 100 897 Skeletonema Genus na 100 2200 5500 30 135 19 294 2 249 100 897 Thlassiosira + Porosira Genus na 30 800 2477 14 597 7893 1500 000 1595 Asterionella + Asterionellopsis Genus 0.03 100 1500 4385 21 815 15 786 1160 000 801 Eucampia + Climacodium Genus 0.03 100	Phytoplankton (cell L^{-1})										
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Pseudo-nitzschia Genus ▲ 0.04 66 1300 6139 47.403 28.064 2.753.000 1300 Paralia Genus na 100 2200 5262 10.539 11.550 919.000 1193 Rhizosolenia Genus ✓ -0.02 100 1600 8000 67.811 44.100 9970.000 1619 Skeletonema Genus na 100 2200 5500 30.135 19.294 2.249.100 897 Thlassiosira + Porosira Genus na 30 800 2477 14.597 7893 1500.000 1595 Leptocylindrus + Tenuicylindrus Genus ▲ 0.07 100 2600 9360 68.055 36.130 11.886.000 879 Asterionella + Asterionellopsis Genus ▲ 0.03 100 1500 4385 21.815 15.786 1160.000 801 Eucampia + Climacodium Genus ▲ 0.03 100 2100 7016 21.829 201.71 349.600 824 Dict	Chaetoceros	Genus	na		10	1900	8700	68 557	36726	5 340 000	1387
Paralia Genus na 100 2200 5262 10 539 11 550 919 000 1193 Rhizosolenia Genus ▼ -0.02 100 1600 8000 67 811 44 100 9970 000 1619 Skeletonema Genus na 100 2200 5500 30 135 19 294 2 249 100 897 Thlassiosira + Porosira Genus na 30 800 2477 14 597 7893 1500 000 1595 Leptocylindrus + Tenuicylindrus Genus A 0.07 100 2600 9360 68 055 36 130 11 886 000 879 Asterionella + Asterionellopsis Genus ▲ 0.07 100 2600 9360 68 055 36 130 11 886 000 879 Asterionella + Asterionellopsis Genus ▲ -0.02 80 800 2450 16 883 8466 2530 000 920 20 Cryptophyceae Class na 100 100 200 578 877 7016 299 Dinophyceae<	Pseudo-nitzschia	Genus		0.04	66	1300	6139	47 403	28 064	2753000	1300
Rhizosolenia Genus ▼ -0.02 100 1600 8000 67 811 44 100 9970 000 1619 Skeletonema Genus na 100 2200 5500 30 135 19 294 2 249 100 897 Thlassiosira + Porosira Genus na 30 800 2477 14 597 7893 1 500 000 1595 Leptocylindrus + Tenuicylindrus Genus △ 0.07 100 2600 9360 68 055 36 130 11 886 000 879 Asterionella + Asterionellopsis Genus △ 0.03 100 1500 4385 21 815 15 786 1160 000 801 Eucampia + Climacodium Genus ▼ -0.02 80 800 2450 16 883 8466 2 530 000 920 Cryptophyceae Class 0.19 100 2100 7016 21 629 20 171 349 600 824 Dictyochophyceae Class 0.04 400 700 2731 8960 8193 2548 000 1538 Prorocentrum	Paralia	Genus	na		100	2200	5262	10 539	11 550	919 000	1193
Skeletonema Genus na 100 2200 5500 30 135 19 294 2 249 100 897 Thlassiosira + Porosira Genus na 30 800 2477 14 597 7893 1 500 000 1595 Leptocylindrus + Tenuicylindrus Genus A 0.07 100 2600 9360 68 055 36 130 11 886 000 879 Asterionella + Asterionellopsis Genus A 0.03 100 1500 4385 21 815 15 786 1 160 000 801 Eucampia + Climacodium Genus Image: Classic intervent of the second intervent of	Rhizosolenia	Genus		-0.02	100	1600	8000	67 811	44 100	9 970 000	1619
Thlassiosira + Porosira Genus na 30 800 2477 14597 7893 1500 000 1595 Leptocylindrus + Tenuicylindrus Genus A 0.07 100 2600 9360 68 055 36 130 11 886 000 879 Asterionella + Asterionellopsis Genus A 0.03 100 1500 4385 21 815 15 786 1160 000 801 Eucampia + Climacodium Genus ✓ -0.02 80 800 2450 16 883 8466 2 530 000 920 Cryptophyceae Class △ 0.19 100 2100 7016 21 629 20 171 349 600 824 Dictyochophyceae Class na 100 100 200 578 877 7016 299 Dinophyceae Class △ 0.03 0.2 200 877 6204 3508 1351 500 540 Prorocentrum Genus △ 0.03 0.2 200 877 6204 3508 1351 500 540 Phaeocysti	Skeletonema	Genus	na		100	2200	5500	30 135	19 294	2 249 100	897
Leptocylindrus + Tenuicylindrus Genus ▲ 0.07 100 2600 9360 68 055 36 130 11 886 000 879 Asterionella + Asterionellopsis Genus ▲ 0.03 100 1500 4385 21 815 15 786 1160 000 801 Eucampia + Climacodium Genus ▼ -0.02 80 800 2450 16 883 8466 2 530 000 920 Cryptophyceae Class ▲ 0.19 100 2100 7016 21 629 20 171 349 600 824 Dictyochophyceae Class na 100 100 200 578 877 7016 299 Dinophyceae Class ▲ 0.03 0.2 200 877 6204 3508 1351 500 540 Phaeocystis globosa Species na 1 6375 731 222 4243 913 4748 154 57734 729 354 Dinophyceae or Bacillariophyceae Ratio △ 0.04 0.0019 0.08 0.02 0.06 0.05 2.49 1517	Thlassiosira + Porosira	Genus	na		30	800	2477	14 597	7893	1 500 000	1595
Asterionellopsis Genus ▲ 0.03 100 1500 4385 21 815 15 786 1 160 000 801 Eucampia + Climacodium Genus ▼ -0.02 80 800 2450 16 883 8466 2 530 000 920 Cryptophyceae Class ▲ 0.19 100 2100 7016 21 629 20 171 349 600 824 Dictyochophyceae Class na 100 100 200 578 877 7016 299 Dinophyceae Class ▲ 0.03 0.2 200 877 6204 3508 1351500 540 Prorocentrum Genus ▲ 0.03 0.2 200 877 6204 3508 1351500 540 Phaeocystis globosa Species na 1 6375 731 222 4243 913 4748 154 57 734 729 354 Dinophyceae or Bacillariophyceae Ratio ▲ 0.04 0.0019 0.008 0.02 0.06 0.05 2.49 1517 Phaeocystis globosa	Leptocylindrus + Tenuicylindrus	Genus		0.07	100	2600	9360	68 055	36 1 30	11 886 000	879
Eucampia + Climacodium Genus Image: Constraint of the system -0.02 80 800 2450 16 883 8466 2 530 000 920 Cryptophyceae Class Image: Class	Asterionella + Asterionellopsis	Genus		0.03	100	1500	4385	21815	15 786	1 160 000	801
Cryptophyceae Class Image: Class Image	Eucampia + Climacodium	Genus		-0.02	80	800	2450	16883	8466	2 530 000	920
Dictyochophyceae Class na 100 100 200 578 877 7016 299 Dinophyceae Class ▲ 0.04 40 700 2731 8960 8193 2548 000 1538 Prorocentrum Genus ▲ 0.03 0.2 200 877 6204 3508 1 351 500 540 Phaeocystis globosa Species na 1 6375 731 222 4243 913 4748 154 57 734 729 354 Dinophyceae or Bacillariophyceae Ratio ▲ 0.04 0.0019 0.08 0.02 0.06 0.05 2.49 1517 Physicochemical <	Cryptophyceae	Class		0.19	100	2100	7016	21 629	20 17 1	349 600	824
Dinophyceae Class Image: Class	Dictyochophyceae	Class	na		100	100	200	578	877	7016	299
Prorocentrum Genus ▲ 0.03 0.2 200 877 6204 3508 1 351 500 540 Phaeocystis globosa Species na 1 6375 731 222 4243 913 4748 154 57 734 729 354 Dinophyceae or Bacillariophyceae Ratio ▲ 0.04 0.0019 0.08 0.02 0.06 0.05 2.49 1517 Physicochemical	Dinophyceae	Class		0.04	40	700	2731	8960	8193	2548000	1538
Phaeocystis globosa Species na 1 6375 731 222 4243 913 4748 154 57 734 729 354 Dinophyceae or Bacillariophyceae Ratio 1 0.004 0.008 0.02 0.06 0.05 2.49 1517 Physicochemical 1	Prorocentrum	Genus		0.03	0.2	200	877	6204	3508	1 351 500	540
Dinophyceae or Bacillariophyceae Ratio 0.04 0.00019 0.008 0.02 0.06 0.05 2.49 1517 Physicochemical	Phaeocystis globosa	Species	na		1	6375	731 222	4 243 913	4 748 154	57 734 729	354
Physicochemical	Dinophyceae or Bacillariophyceae	Ratio		0.04	0.00019	0.008	0.02	0.06	0.05	2.49	1517
•	Physicochemical										
Temperature (°C) 1 .80 1 .80 1 .80 1 .60 1 .95 1 7.30 1 .30 2 .20 2 .292	Temperature (°C)			0.003	1.80	8.90	12.60	12.95	17.30	23.20	2292
Salinity na 31.40 33.60 33.90 33.87 34.20 37.16 913	Salinity		na		31.40	33.60	33.90	33.87	34.20	37.16	913
NH ₄ (μ molL ⁻¹) na 0.03 2.42 4.79 5.32 7.21 49.00 1509	$NH_4 \ (\mu mol L^{-1})$		na		0.03	2.42	4.79	5.32	7.21	49.00	1509
NO ₂ (μ molL ⁻¹) na 0.01 0.23 0.40 0.51 0.65 16.00 902	NO ₂ (μ mol L ⁻¹)		na		0.01	0.23	0.40	0.51	0.65	16.00	902
NO ₃ $(\mu mol L^{-1})$ na 0.09 1.61 4.40 8.73 14.90 57.00 1310	NO ₃ (μ mol L ⁻¹)		na		0.09	1.61	4.40	8.73	14.90	57.00	1310
$NO_2 + NO_3$ (umol L ⁻¹) na 0.09 1.87 4.69 9.02 15.16 57.70 1099	$NO_2 + NO_3 (\mu mol L^{-1})$		na		0.09	1.87	4.69	9.02	15.16	57.70	1099
$PO_1(umolL^{-1})$ $-0.01 0.05 0.32 0.58 0.79 1.05 1.05 2.89$	PO_4 (umol L^{-1})		•	-0.01	0.05	0.32	0.58	0.79	1.05	1.05	289
SiOH (umol L^{-1}) na 0.1 2.00 4.55 5.27 8.01 21.51 242	SiOH (umol L^{-1})		na		0.1	2.00	4.55	5.27	8.01	21.51	242
Chlorophyll a (ugL ⁻¹) -0.02 0.01 1.38 2.85 4.90 6.26 60.76 1901	Chlorophyll a (ugL ⁻¹)			-0.02	0.01	1.38	2.85	4.90	6.26	60.76	1901

4 Database

To efficiently manage coastal monitoring data, IFREMER has developed the Quadrige² information system (https: //envlit.ifremer.fr/Quadrige-la-base-de-donnees, last access: 12 April 2023), which combines a database with various products and services. Quadrige² plays a crucial role in two key areas: (1) securely and optimally storing basic monitoring data, including analysis results from all the monitoring networks, in a supervised and scalable manner; and

(2) interpreting and enhancing the value of these data. Once the data are stored with an assigned quality level, they become available for a wide range of applications. As a result, this system is the required link for monitoring data between data collection in the field and their availability in multiple formats. Quadrige² has been approved as the national reference information system for coastal waters by the French Ministry of the Environment and is part of broader national data portals dedicated to ocean data such as ODATIS



Figure 2. Time series of the different physicochemical parameters and chlorophyll-*a* concentrations measured from 1978 to 2023 as part of the IGA Gravelines monitoring program at the Canal d'amenée sampling station.

Pole, which is part of the research e-infrastructure Data Terra (https://www.data-terra.org/, last access: 12 June 2024).

The datasets presented in this article are derived from extractions from the Quadrige² database. The raw extraction, which includes all the parameters (hydrological, phytoplankton, and zooplankton), is available from SEANOE. The data are provided as a semicolon-delimited .csv file, but they contain non-ASCII characters and a few errors that have accumulated over time. Consequently, a data preprocessing phase was applied to the extraction, including checks for duplicates, outliers, the accuracy of the identified taxa, and the verification and harmonization of the measurement units. This ensured that the dataset was clean and reliable for further analysis.

Some cells in these databases are empty, particularly in the "Hours" column, where the sampling time was not consistently recorded by operators. Additionally, the metadata for the "NO₃ + NO₂" parameter are only filled when the values result from direct sample analyses, not from the addition of NO₂ and NO₃ results, which was done for database harmonization. Consequently, the final quality-controlled database is divided into three separate semicolon-delimited .csv files, all encoded in UTF-8 with ASCII characters and using a dot as the decimal separator. The first file, containing hydrochemical parameters and chlorophyll-*a* concentrations, has

19 columns and 12 635 rows. The second file, which holds phytoplankton abundance data, has 20 columns and 37 418 rows. The third file, dedicated to zooplankton abundance data, has 21 columns and 12 453 rows.

The database header columns are in French, as Quadrige² is a French national database. A French–English translation of these headers is provided in the Supplement, and a detailed description can be found in IFREMER (2017). Physicochemical, phytoplankton, and zooplankton data are available for the 1978–2023 period, because the database was up to date until 2023 at the time of writing. The datasets will be updated annually in SEANOE, maintaining the same DOI to ensure continuity and accessibility.

5 Quality control

5.1 Data validation

The data are collected in the field and/or laboratory and subsequently entered into the Quadrige² database via a dedicated user interface. Data control involves verifying and potentially modifying the entered data (including both results and metadata) in order to ensure consistency with the original bench book (or field sheets). After this verification and any necessary corrections, the data follow this validation process:



Figure 3. Annual box–whisker plots of the main physicochemical parameters at the Canal d'amenée sampling station as part of the IGA Gravelines monitoring program over the period 1978–2023. For improved data visualization, outliers are not represented.

- Data validity this ensures the accuracy and reliability of the data corresponding to the analytical results.
- Data locking this secures the data, preventing further modifications, even by the original data entry person.
- Data distribution once verified, the data become accessible for extraction and dissemination by all authorized Quadrige² users.

5.2 Data qualification

Following the initial round of data verification, the data undergo a qualification procedure that involves the following:

- Search for data that may be scientifically suspicious or clearly erroneous or aberrant.
- Correct data where possible, making adjustments to correct any identified issues.
- Assign a qualification level as follows.
 - Good: the data are scientifically valid and relevant.
 - Doubtful: the data may be inaccurate, and taking them into account may bias the results.

 False: the data are considered erroneous or problematic and should not be included in the analysis.

The level of qualification reflects the level of confidence in the data. It determines the way in which the data are distributed (only data qualified as "good" and "doubtful" are widely distributed) and how they are used in specific data processing. This is determined through a two-step qualification process:

- 1. "automatic" qualification, which involves identifying obvious and easily detectable errors in the data; and
- "expert" qualification, which focuses on detecting statistically aberrant data using adapted methods (e.g., time series analysis or statistical tests). Only data qualifying as good or doubtful from the previous step are used for this expert qualification.

6 Data analysis

The R package TTAinterfaceTrendAnalysis was used to efficiently, homogeneously, and rapidly perform these tests and extract the most relevant statistical metrics (Devreker and Lefebvre, 2014). Temporal trend analysis was performed using the seasonal Mann–Kendall nonparametric test, with p values corrected for autocorrelated data. Since this test



Figure 4. Monthly box–whisker plots of the main physicochemical parameters at the Canal d'amenée sampling station as part of the IGA Gravelines monitoring program over the period 1978–2023.

is suited for monotonic trend analyses, the cumulative sum method was applied to identify shifts within the time series. Moreover, when a significant trend was detected, the Theil– Sen slope estimator was used to quantify the magnitude of the trend.

7 Data summary

Table 3 presents the descriptive statistics for each physicochemical and biological parameter at the Canal d'amenée station. At an interannual scale, temperature is the only hydrological parameter showing a significant upward trend between 1978 and 2023, with an average increase of +0.04 °C yr⁻¹, resulting in a total rise of +1.8 °C during the survey period. Among nutrients, only phosphorus shows a significant decreasing trend, with a rate of $-0.01 \,\mu\text{mol}\,\text{L}^{-1}\,\text{yr}^{-1}$. Chlorophyll-*a* concentrations also demonstrate a significant decreasing trend with a clear shift in the time series (Figs. 2 and 3 and Table 3). This shift, detected by the Pettitt test, occurs around 2012, with mean chlorophyll-*a* concentrations decreasing from $5.85 \,\mu g L^{-1}$ before 2012 to $2.68 \,\mu g \, L^{-1}$ afterwards (and from 8 to $3.7 \,\mu g \, L^{-1}$ during the growing season from March to September).

At the inter-seasonal scale, nutrient concentrations (nitrate + nitrite, silicate, and phosphorus) are maximal in winter (Fig. 4). These concentrations decrease in February and March, when concentrations in chlorophyll *a* (Fig. 4) and phytoplankton abundances (Fig. 5) increase. Chlorophyll*a* concentrations reach their maximum from March to July, with the highest average value occurring in April $(12 \,\mu g L^{-1})$ and the maximum value reached in March 2010 at > $60 \,\mu g L^{-1}$. Temperature follows a typical pattern for temperate coastal waters, with February being the coldest month ($6.9 \,^{\circ}$ C) and August the warmest ($19.6 \,^{\circ}$ C) (Fig. 4). Mean oxygen concentrations are highest in winter (~ 10 to $11 \,\text{mg} L^{-1}$), when photosynthetic activity and temperature are lowest. These levels decrease in spring, reaching 8 to $9 \,\text{mg} L^{-1}$ (Fig. 4).

Interannual variability of phytoplankton communities reveals a significant increase in the abundance of some taxonomic classes, including Dinophyceae and Cryptophyceae. The species *Phaeocystis globosa* does not show any significant trend when some Bacillariophyceae show significant increasing trends. On an annual scale, phytoplankton communities are typically dominated by either *P. globosa* or Bacillariophyceae, alternating between the two (with Bacillariophyceae dominance observed in 1993–1997, 2002–2005, and 2013) (Fig. 6). Mean annual abundances tend to be higher



Figure 5. Weekly average variability (1978–2023) of the major phytoplankton groups (Prymnesiophyceae, mainly *Phaeocystis globosa*, Bacillariophyceae, Dinophyceae, Cryptophyceae, and other phytoplankton) at the Canal d'amenée sampling station, as part of the IGA Gravelines monitoring program. The vertical bars illustrate the relative abundances of these groups (%), while the black circles indicate the mean weekly total phytoplankton abundance (10^5 cells L⁻¹).

during years when *P. globosa* is dominant (Fig. 6). The Dinophyceae / Bacillariophyceae abundance ratio (an indicator of eutrophication status; Wasmund et al., 2017; Xiao et al., 2018) shows a significant increasing trend, suggesting that Dinophyceae abundances are increasing more rapidly than Bacillariophyceae abundances (Table 3).

Phytoplankton communities also show a clear seasonal pattern. Abundances are low in early winter and begin to increase with Bacillariophyceae starting in February. From March to May, *Phaeocystis globosa* is dominant, constituting over 90% of the total phytoplankton abundance, while Bacillariophyceae remain dominant for the rest of the year. *Phaeocystis globosa* bloom corresponds to the peak period of phytoplankton abundance and the lowest specific richness (Fig. 7). During the average *P. globosa* bloom, Bacillariophyceae species such as *Chaetoceros* or *Asterionellopsis* are present at the beginning, followed later by *Guinardia* and *Pseudo-nitzschia*.

Concerning zooplankton communities, the mean seasonal variability shows dominance of meroplankton in February, while copepods are clearly dominant during the rest of the year (Fig. 8). Zooplankton mean monthly abundance shows a peak value in May of 7200 ind m⁻³ when the copepod *Temora longicornis* is the dominant species (Fig. 8), which corresponds to a slight decrease in zooplankton-specific richness (Fig. 7). Copepod succession shows large dominance

of calanoids (*T. longicornis, Acartia clausii, Centropages, Paracalanus*, and *Pseudocalanus*) from January to July, and harpacticoid relative abundance increases in August to represent 50% of the zooplankton community in September (Fig. 9).

There is no clear interannual trend in overall zooplankton community abundance (Table 3). However, meroplankton abundance shows a significant decreasing trend over time. Among copepods, *Paracalanus* shows a significant increasing trend, while *Pseudocalanus* shows a significant decreasing trend. Mean interannual zooplankton abundances demonstrate some variability, with copepod species remaining the dominant group (Fig. 10).

The diversity of both zooplankton and phytoplankton communities follows similar seasonal patterns, with the lowest diversity in April and May during the *P. globosa* bloom (Fig. 7). However, there is a notable difference in February and March, when phytoplankton diversity increases while zooplankton diversity decreases.

8 Code and data availability

The IGA-HP Gravelines dataset is publicly available at https://doi.org/10.17882/102656 (Lefebvre et al., 2024) (hydrology and plankton monitoring program at the Gravelines coastal station in 2024).



Figure 6. Interannual variability of the major phytoplankton groups (Prymnesiophyceae, mainly *Phaeocystis globosa*, Bacillariophyceae, Dinophyceae, Cryptophyceae, and other phytoplankton) at the Canal d'amenée sampling station, as part of the IGA Gravelines monitoring program. The vertical bars represent the relative abundances of these groups (%), while the black circles indicate the mean annual total phytoplankton abundance $(10^5 \text{ cells L}^{-1})$.



Figure 7. Mean seasonal variation (monthly scale, 1978–2023) of the Shannon–Wiener diversity index for both phytoplankton and holo(zoo)plankton at the Canal d'amenée sampling station, as part of the IGA Gravelines monitoring program.

The R package TTAinterfaceTrendAnalysis is available on the CRAN website (Comprehensive R Archive Network – https://cran.r-project.org/package= TTAinterfaceTrendAnalysis/index.html, Devreker and Lefebvre, 2021).

9 Discussion and conclusion

The IGA Gravelines Canal d'amenée data series, which began in 1978, represents the longest physicochemical and planktonic data series in the South Bight of the North Sea. This time series captures the dynamics of phytoplankton, mainly dominated by Bacillariophyceae and Prymne-



Figure 8. Mean seasonal variability (1978–2023) of the major zooplankton groups (Appendicularia, Branchiopoda, Copepoda, Chaetognatha, Ctenophora, meroplankton, and other holoplankton) at the Canal d'amenée sampling station of the IGA Gravelines monitoring program. The vertical bars indicate the relative abundances of these groups (%), while the black circles represent the mean monthly total abundance $(10^2 \text{ ind m}^{-3})$.



Figure 9. Mean monthly variability (1978–2023) of copepod orders (Calanoida, Cyclopoida, and Harpacticoida) at the Canal d'amenée sampling station of the IGA Gravelines monitoring program. The vertical bars illustrate the relative abundances of these groups (%), while the black circles show the mean monthly total abundance $(10^2 \text{ ind m}^{-3})$.

siophyceae (*Phaeocystis globosa*) as well as zooplankton species, with a focus on copepods. The dataset reveals temporal successions and multiyear trends in plankton communities, including annual phenology characterized by an initial increase in Bacillariophyceae abundance, followed by the early spring bloom of *P. globosa*, which coincides with high abundances of the potential harmful alga *Pseudonitzschia* sp.

The Bacillariophyceae and *P. globosa* bloom is one of the most structuring events in the planktonic community of this region of the eastern English Channel (Schapira et al., 2008; Grattepanche et al., 2011), influencing not only plankton, but also bacteria (Lamy et al., 2006) and benthic communities (Dauvin et al., 2008; Denis and Desroy, 2008; Spilmont et al., 2009). The IGA survey shows that this bloom varies significantly in intensity and relative proportion be-



Figure 10. Interannual variability of the major zooplankton groups (Appendicularia, Branchiopoda, Copepoda, Chaetognatha, Ctenophora, meroplankton, and other holoplankton) at the Canal d'amenée sampling station of the IGA Gravelines monitoring program. The vertical bars indicate the relative abundances of these groups (%), while the black circles represent the mean annual total abundance $(10^2 \text{ ind m}^{-3})$.

tween years. During this bloom, annual chlorophyll-*a* concentrations peak, while nutrient levels hardly decrease. This period also sees a significant drop in the annual proportion of copepod species (the most abundant primary consumers in the zooplankton community), showing the impact of this bloom on the entire pelagic ecosystem, as previously described in this region (Lancelot et al., 2005).

The interannual variability in plankton dynamics was concomitant with a significant increasing trend in seawater temperature and a significant decreasing trend in phosphate concentrations. Both of these parameters were already described as drivers of *P. globosa* blooms (Hernández Fariñas et al., 2015; Karasiewicz et al., 2018). Given the variety of the parameters measured, the frequency of the measurements, and the length of the dataset, this series is exceptionally well suited to studying the dynamics of planktonic communities and the effects of anthropogenic pressures on them.

Indeed, similar long-term data series have proven to be valuable resources for scientific research, leading to a significant number of publications across various topics. For instance, the Point B series from the Laboratoire d'Océanographie de Villefranche (LOV) on the Mediterranean Sea (Romagnan et al., 2015, 2016; Feuilloley et al., 2022), the L4 series from the Plymouth Marine Laboratory in the western English Channel (Harris, 2010; Widdicombe et al., 2010; McEvoy et al., 2023), the SRN series from IFRE-MER's LER-BL laboratory in the eastern English Channel and North Sea (Lefebvre and Devreker, 2023; Lefebvre et al., 2011), and other extensive pan-European series such as the Continuous Plankton Record survey (Holland et al., 2024) or those reporting to the OSPAR biodiversity working group (Holland et al., 2023) have generated a substantial number of scientific articles addressing a wide range of topics. These include plankton dynamics (Vandromme et al., 2011; John et al., 2001; Lefebvre and Devreker, 2023), climate change (Corona et al., 2024; Kapsenberg et al., 2017; Parravicini et al., 2015), HABs (Karasiewicz et al., 2020; Karasiewicz and Lefebvre, 2022), food webs (Atkinson et al., 2015), and the carbon cycle (González-Benítez et al., 2019). Such extensive, long-term, and multiparameter datasets can also be critical for assessing environmental quality within the frameworks of European directives (such as the Water Framework Directive – WFD – or the Marine Strategy Framework Directive - MSFD) and regional marine policies (such as the OSPAR or Barcelona conventions) (McQuatters-Gollop et al., 2019; Lefebvre and Devreker, 2020).

The IGA Gravelines Canal d'amenée series has been used in various studies, including analyses of long-term temperature fluctuations and their link with the North Atlantic Oscillation (NAO) (Woehrlings et al., 2005). The dataset, covering measurements from 1975 to 1992, has been used to investigate trophic relationships (Le Fevre-Lehoerff et al., 1993) and cycles within the context of climate changes (Le Fevre-Lehoerff et al., 1995), incorporating data on water temperature, salinity, suspended matter, nutrients, chlorophyll a, and zooplankton. Halsband-Lenk and Antajan (2010) showed the utility of this time series for defining regional multi-metric food web indices using temperature, salinity, chlorophyll pigments, and phytoplankton and zooplankton abundance data. The series has also been identified as one of interest by the ICES WGZE (Working Group on Zooplankton Ecology) and has been integrated into NOAA's METABASE (https: //www.st.nmfs.noaa.gov/copepod/time-series/fr-30101/, last access: 20 September 2024). Despite the highest scientific value of this dataset, the number of publications focused on plankton and hydrology remains relatively limited compared to other datasets, such as L4 and Point B (Google Scholar results for "plankton" and the station name since 2020: 128 for L4 and 69 for Point B). Publications related to benthic fauna from the IGA Gravelines survey are more numerous.

Moreover, studying plankton dynamics using such data series can provide critical insights into preventing cooling problems at nuclear power plants. High biomass blooms can obstruct cooling systems by either physically blocking the flow or altering water viscosity, leading to reduced efficiency and potential operational issues. Such events have been documented in the literature, including cases involving gelatinous species at Gravelines (Antajan et al., 2014) and HABs globally (Wang et al., 2022). At the GNPP, in addition to issues caused by gelatinous blooms, particularly from Pleurobrachia pileus, blooms of Phaeocystis globosa can also affect cooling systems. Consequently, it is crucial to understand the drivers of P. globosa blooms, including the parameters that influence their intensity and timing. Developing early warning systems to anticipate such blooms and implementing preventive or mitigative measures is thus essential for maintaining operational efficiency and avoiding disruptions.

The data acquired through the IGA program follow findability, accessibility, interoperability, and reusability (FAIR) principles, which define a set of rules to facilitate FAIR principles of data and the associated metadata. This dataset from Gravelines is particularly well suited to collaborative annotation work, which could enhance forecasting capabilities. This process involves identifying and labeling recurring, rare, and extreme events in the time series by experts. Recent advancements in machine learning have introduced several tools to assist with pattern recognition and automatic segmentation of time series data. Techniques include fixedlength window segmentation (Van Hoan et al., 2017), sliding window approaches using autoencoders (Längkvist et al., 2014), expectation-maximization models (Poisson-Caillault and Lefebvre, 2017), hidden Markov models (Dias et al., 2015; Rousseeuw et al., 2015), and multilevel spectral clustering (Grassi et al., 2020). These methods can be used to isolate patterns in the time series, which can then be annotated with relevant event labels based on the expertise of the annotators.

This labeled database could be used as a reference training set by the scientific community, addressing specific needs in artificial intelligence. It would be particularly valuable for developing algorithms, calibrating models, and implementing digital prediction and warning systems. Such advancements could enhance biologists' understanding of marine dynamics, providing new insights into plankton community functioning and environmental state evolution, including potential trends or regime shifts in the context of global environmental change.

The recent decision by EDF (Électricité de France) to provide open access to the IGA Gravelines Canal d'amenée monitoring data offers significant opportunities for scientific advancement. This will facilitate the utilization of the time series in ways similar to other long-term datasets. By detailing the characteristics of the Gravelines Canal d'amenée series, this article provides researchers and managers with the information needed for future scientific investigations and management applications, thereby contributing to a deeper understanding of coastal ecosystems in the English Channel and North Sea.

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