



The Western Channel Observatory: a century of oceanographic, chemical and biological data compiled from pelagic and benthic habitats in the Western English Channel.

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Abstract

The Western Channel Observatory (WCO) comprises a series of pelagic, benthic and atmospheric sampling sites within 40

- 20 km of Plymouth UK, which have been sampled by the Plymouth Institutes on a regular basis since 1903. This longevity of recording and the high frequency of observations provide a unique combination of data; for example temperature data were first collected in 1903 and the reference station L4 has been sampled on a weekly basis since 1988 where nearly 400 planktonic taxa have been enumerated. While the component datasets have been archived, here we provide the first summary database bringing together a wide suite of the observations. This provides monthly average values of some of the key pelagic
- and benthic measurements for the inshore site L4 (50° 15.00' N, 4° 13.02' W approx depth 55 m), the offshore site E1 (50° 02.00' N, 4° 22.00' W approx depth 75 m) and the intermediate L5 site (50° 10.80' N 4° 18.00' W approx depth 58m. In brief, the data include: water temperature (from 1903); macronutrients (from 1934); dissolved inorganic carbon and total alkalinity (from 2008); methane and nitrous oxide (from 2011); chlorophyll *a* (from 1992); HPLC-derived pigments (from 1999); <20µm plankton by flow cytometry including bacteria (8 functional groups from 2007); phytoplankton by microscopy (6</p>
- 30 functional groups from 1992); microplankton and mesozooplankton from FlowCam (6 groups from 2012), *Noctiluca* sp. dinoflagellate (from 1997); mesozooplankton by microscopy (8 groups from 1988); *Calanus helgolandicus* egg production rates (from 1992); fish larvae from Young Fish Trawl survey (4 groups from 1924); benthic macrofauna (4 groups from 2008); demersal fish (19 families from 2008); blue shark, *Prionace glauca* (from 1958); 16S alpha diversity for sediment



and water column (from 2012). These data have varying coverage in time and depth resolution. The metadata tables describe 35 each data set, provide pointers to the source data and other related Western Channel Observatory data sets and outputs not compiled here. We provide summaries of the main trends in seasonality and some major, climate related shifts that have been revealed over the last century. The data are available from Data Archive for Seabed Species and Habitats (DASSH) via the link http://doi.org/10.17031/645110fb81749 (McEvoy and Atkinson, 2023). Making the data fully accessible and including units of both abundance and biomass will stimulate a variety of uptakes. These may include uses as an educational resource for projects, for models and budgets or for analysis of seasonality and long-term change in a coupled benthic-40 pelagic system and for supporting UK and Northeast Atlantic policy and management.

1 History

Sustained observations of the marine environment are vital to understand marine ecosystem functioning and climate change responses (O'Brien et al., 2017; Richardson, 2008). Over seasonal timescales, high resolution observations allow the understanding of community succession and seasonality (Smyth et al., 2014) and over multiple decades they allow us to 45 tease out the effects of local variability and anthropogenic stressors from the longer-term signal of climate change (Edwards & Richardson, 2004; Ratnarajah et al., 2023). Paradoxically, however, many sampling programs are funded for only 3-4 years and despite the urgency of understanding climate change responses, time series globally are threatened (Vucetich et al., 2020) This makes it even more important to make data from existing long time series findable, available for re-use and as 50 well documented as possible.

The Western Channel Observatory (WCO) data contains an unprecedented collection of parameters both in terms of longevity and variety. Investigation of the marine environment in the western English Channel off Plymouth began with the opening of the Marine Biological Association (MBA) laboratory in 1888. Given the importance in the area of the pelagic

- 55 fishery the remit focused strongly on research in applied fisheries. Initial studies centered on the eggs and larvae of commercially important fish. With the advent of the International Council for the Exploration of the Sea (ICES) and a growing realisation that hydrography had an influence on biological communities, plankton surveys and hydrographical measurements were soon added (Southward et al., 2005; Southward & Roberts, 1987). In the decades that followed, observations were expanded with the creation of stations E1(50° 02.00' N, 4° 22.00' W) and L5(50° 10.80' N 4° 18.00'
- 60 W). Sampling was interrupted during both World War I (1914-1918) and World War II (1939-1945). Funding priorities and organisational changes in the 1970s and 1980s threatened the future of long-term time series, and sampling at L5 and E1 was consequently stopped until 2002. However, in 1988 Plymouth Marine Laboratory (PML) established weekly zooplankton sampling at station L4(50° 15.00' N, 4° 13.02' W), with ad-hoc funding and no formal support. Sampling for phytoplankton community composition and abundance egg production and environmental variables followed from 1992 onwards. The
- 65 WCO was founded in 2005 to bring these valuable time series together. The WCO provided a platform for a wider array of





parameters to be initiated, for example the benthic survey from 2007, *in-situ* automated buoys at L4 and E1 (supported initially by Natural Environmental Research Council (NERC) and then the Met Office), the Penlee Point Atmospheric Observatory (PPAO) from 2014 and Smart Sound Plymouth from 2021 (**Fig 1**).

- 70 The stations around Plymouth now known as the Western Channel Observatory have supported major innovative work, for example pioneering work on plankton as indicators (Russell, 1935), the measurement of nutrients and primary production (Boalch et al., 1978), early work on fatty acids and the importance of food quality for zooplankton (Conover & Corner, 1968; Pond et al., 1996) and the use of molecular biology tools to provide insight into the seasonal dynamics of viral and bacterial plankton (Lindeque, 2023; Gilbert et al., 2009; Schroeder et al., 2003). These works, including the development of
- 75 intertidal research and data not covered here can be found in the historical review of Southward et al. (2005). Later Special journal issues cover the 20th and 25th anniversaries of regular sampling at L4 and are described respectively in (Harris, 2010) and (Smyth et al., 2015). We refer the reader to these for the historical context of the observations we summarise here.



Figure 1: Location of Western Channel Observatory (WCO) sampling stations



2 The WCO environment

The two main marine stations of L4 and E1 both exhibit strong seasonal signals and are tidally influenced (Smyth et al., 2015). Both become stratified typically after April, continuing through the summer months and lasting until late September. Station L4 is classified as a coastal site and is periodically influenced by flood water discharge from the rivers Tamar and Plym (Rees et al., 2009). However, at a depth of approx. 55 m and 13 km offshore it is not as prone to localized inshore effects and is classified as "transitionally stratified" (Pingree, 1980). The deeper station E1, 40 km offshore and approx. 75 m deep, is less influenced by coastal water influx and is classified as an open shelf station that is seasonally stratified. The intermediate station, L5, was much sampled in early years and is just west of Eddystone reef. These stations experience classic, albeit highly variable seasonal production cycles with spring and autumn phytoplankton blooms. Figure 2 compares the key WCO sites in relation to the wider summer pattern of stratification (**Fig. 2a**) and to the longer trend of climatic

2b) the key web sites in relation to the wheel summer pattern of straine atom (Fig. 2a) and to the longer tiend of enhance cycles across the North Atlantic (Bode, 2023) highlighting a recent phase of intense warming over the last 4 decades (Fig. 2b). These environmental changes, and the response of the biota, are described more fully in Section 5 using plots derived from our summary database.

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Figure 2: Wider-scale spatial and temporal context for the Western Channel Observatory. (a) The wider setting of the L4 and E1 stations in the Western English Channel, in relation to summer sea surface temperature. Cold colours represent tidally mixed areas, warm colours represent summer stratifying areas with seasonal summer thermocline. L4 stratifies in summer and is defined as transitionally stratified, whereas E1 is open shelf and defined as seasonally stratified (Pingree, 1980). (b) Annual surface

- 125 temperature records at station E1 spanning 1903-2021. Due to missing data in some months of the early years, annual means were calculated here as averages of February, May, August and November. Missing months were interpolated as mean respective month over the whole timespan. Years with more than 2 of the four missing months are not plotted here. Dotted line is least squares linear regression over the whole timespan. (Southward et al., 2005) based on data from last century denoted three thermal 130
- epochs coloured here. Warming was from 1921-1961, followed by a cooling era from 1962-1985 and then a warming period from 1986 to present.



3 Objectives

The individual data sets of the WCO are valuable, but differing levels of reporting and formatting hamper their use and prevent integration. Many are currently available through data repositories such as British Oceanographic Data Centre (BODC) and Data Archive for Seabed Species and Habitats (DASSH), however, some are lodged with individual scientists. To improve their overall utility, the various component data sets need to be brought together into a single format. We have done this here for the first time, but to make this project tractable we have summarized the core datasets as monthly averages, and for broad functional groups. This level of resolution (coarser than some of the measurements, which can be weekly and for individual species), was chosen as a first step to allow timely completion of this initiative, to provide a summary database that combines many diverse data sources. This data paper combines in a single spreadsheet most of the key variables which have good seasonal or longer-term coverage (**Table 1**). Specialists who wish to access the underlying

sets not summarized here are directed to our WCO data catalogue: <u>https://www.westernchannelobservatory.org.uk/data.php.</u>
This catalogue provides sampling details, doi's of the most recent versions, and points of contact for specific data sets.
Additional information is also available in **Table A1** and **Table A2**.

high-resolution observations, data for individual species, who require the most recent data available, or require other data

This data paper is aimed towards scientists who may not need weekly resolution or species-specific data, but who wish to compare the monthly-averaged physical, chemical and biological data. Biological data are provided in units of both abundance and biomass, to enhance their utility for modelling. We have also made the spreadsheets as user-friendly and simple as possible to be of help as an educational resource at the undergraduate level. This data paper describes the database

(Section 4); illustrates its utility to examine seasonality and longer-term trends while summarizing previous work on these topics (Section 5); provides a broader-scale context for the WCO (Section 6) and finally provides practical advice on the strengths, limitations and how to use and cite this database (Sections 7 and 8).

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Table 1: Data combined in this data paper, showing the timespan of each.

4 Data processing

- 180 This paper consolidates 22 individual and diverse data sets using monthly averages. Data with comprehensive seasonal coverage which span at least two years are included. Detailed information on sampling and analysis protocols plus data coverage can be found in the Appendices (**Table A1 and A2**). It is essential to read these appendices before extracting data to avoid errors, for example in distinguishing between zeros and absent data. A zero represents a parameter that was either looked for and not found (for plankton data) or was below the detection limit (nutrient data). A blank cell, by contrast, is
- 185 where there are no data available for that particular month. To benefit models, budgets and size-based approaches, biotic data are reported both in units of abundance and biomass. The only exception is the smallest plankton measured by flow cytometry. These use fixed conversion factors for the whole functional group, and have multiple groups and depths. Therefore to remove the complexity of having many data fields that are simple multiples of others, these are reported only as abundance per millilitre. Median cell diameters are provided, which enables estimation of biomass based on the volume of a
- 190 sphere and carbon values from the literature (**Table 2**). Median cell diameters were derived by collecting seawater samples, filtering them sequentially through a series of membrane filters, analysing the filtrates by flow cytometry and the percentage of cells remaining plotted as a percentage of unfiltered seawater against filter pore size (Burkill et al., 1993).



195 Table 2: Median cell diameters for plankton groups quantified by flow cytometry. [§]from Station L4, approximately monthly over an annual cycle 2013-2014 (unpublished); [^] from the Celtic Sea, April 2002 (unpublished). [£] (Heywood et al., 2006). *Carbon conversion factor 0.22 pg C per µm³ (Booth, 1988), [#]carbon conversion factor 0.285 pg C per µm³ (unpublished).

Group	Synechococcus sp. ^{\$}	Picoeukaryotes ^{\$}	Nanoeukaryotes	Cocco-lithophores^	Crypto-phytes ^{\$}	Bacteria [£]
Median diameter (µm) ±1 SD	1.72 ± 0.70	1.83 ± 0.58	5.40 ± 2.04	7.68 ± 0.89	5.48 ± 1.33	-
Spherical volume (µm ³)	2.66	3.20	82.50	236.87	86.36	-
Carbon per cell (pg)	0.59*	0.70*	18.15*	67.51#	19.00*	0.019

5 Results and Discussion

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In this section we briefly showcase some of the key data sets, by outlining the seasonality and environmental variability, illustrating the coverage of all of the component data series at L4 (Fig. 3) and E1 and L5 (Fig. 4). We then show selected examples of the time series data coverage, including longer term trends at L4 (Fig. 5) and E1 and L5 (Fig. 6). A few other key example results are shown, including the *Calanus* egg production experiments (Fig. 7) and the time-depth resolution of sampling for carbonate chemistry (Fig. 8).

5.1 Overall seasonality: L4

- 205 The high resolution of sampling multiple parameters at L4 makes it an ideal test site for improving understanding of the drivers of seasonality. Based on direct empirical observations and from models derived from them, varying alternative mechanisms have been proposed to drive plankton seasonality (Atkinson et al., 2018) Bentho-pelagic connectivity has been found to be seasonally variable in terms of the origin of the suspended and dissolved matter fluctuating between the two ecospheres, as well as the dominant flux directions (Queirós et al., 2015; Rühl et al., 2020; Tait et al., 2015). Figure 3
- summarises some of the key aspects of this seasonality at L4. In brief, L4 is a transitionally stratified site (Pingree, 1980) that 210 stratifies typically from around May to September with surface temperatures ranging from about 9°C in March to around 16°C in August. There is, however, strong biological connectivity between surface waters and the seabed at L4, even during stratified periods, as illustrated by pigment data and stable isotopic signatures of both dissolved inorganic carbon (DIC) and particulate organic carbon and nitrogen (Queirós et al., 2019; Tait et al., 2015). This stratification cycle drives much of the
- 215 pelagic dynamics with nutrient (especially nitrate) depletion to near-limiting levels at the limit of detection in the upper water column during the stratified period, as well as progressive reductions in DIC, Methane and Nitrous Oxide typically until about August. (Kitidis et al., 2012).





The combination of nutrients, light and grazing cause the conditions for a "classic" temperate shelf sea production cycle (Irigoien et al., 2005; Kiørboe & Hirst, 2008) with varying interpretations of causal mechanisms as described above. Thus there is typically a spring bloom around April-May dominated by diatoms and the prymnesiophyte *Phaocystis*, followed by a dinoflagellate bloom in late summer and often diatoms in the autumn. Importantly, however, the averages in **Fig. 3** disguise substantial inter-annual variability (Widdicombe et al., 2010) with sometimes a single summer bloom. The pico- and nano fractions follow slightly different dynamics, with highest biomasses building up in the summer stratified period with maxima

- 225 often in August-September (Tarran & Bruun, 2015). At the size boundary between phytoplankton and metazoans, the Lugol's based counts are imprecise due to rarity, yet the organisms are too small to be quantitatively retained by the 200 μm mesozooplankton WP2 nets. In this size range the FlowCam biomass estimates based on 63 μm mesh, full-depth net hauls show important contributions of copepod nauplii and the larger diatoms, dinoflagellates and ciliates. Rare seasonal profile data of these larger forms in relation to the copepod *Oithona* life stages based on bottle sampling is provided by (Cornwell et
- 230 al., 2020).

The mesozooplankton grazers from the full depth 200 μ m net hauls follow slightly counter-intuitive dynamics. Paradoxically, they tend to increase substantially as early as March, pre-empting the spring bloom and grazing the early start of the diatom bloom (Atkinson et al., 2015). The peak is typically in the early summer, dominated both numerically (Eloire et al., 2010) and in terms of biomass by copepods, but also having a substantial contribution in spring from meroplankton

- (Highfield et al., 2010). More predatory taxa (often gelatinous or semi-gelatinous forms such as chaetognaths) then become important later in summer. Egg production rate of *Calanus helgolandicus* has for most of the time series been highest in the April-June spring bloom months (Irigoien & Harris, 2003; Maud et al., 2015; Maud et al., 2018) although as described in section 5.5 this is changing. This copepod species alongside other zooplankton such as appendicularians (López-Urrutia et al., 2003) decapods (Fileman et al., 2014), bivalve larvae (Lindeque et al., 2015) and *Oithona similis* (Castellani et al., 2014).
- 2016; Cornwell et al., 2018; Cornwell et al., 2020) has been the focus of a series of detailed studies at L4 (Bonnet et al., 2005; Hirst et al., 2007; Irigoien & Harris, 2003).

In contrast to the plankton, the benthic and demersal taxa have more varied seasonal dynamics. Macrofauna biomass is dominated at L4 by suspension feeders with similar biomasses for most of the year (**Fig. 3**) except for depressed values in early winter. Potentially reflecting seasonal variation in water column food supply, species richness of infauna peaks

- 245 throughout the summer in surface sediments and is lowest in late Autumn. Higher numbers of species are also found in deeper sediment layers during warmer months, with the community seemingly shallowing over winter (Queirós et al., 2019). An assessment of particulate carbon sources to the seabed at L4 also suggested that fauna in shallow sediment layers exhibit strong signals of suspension and deposit feeding reliant on planktonic food sources, with carnivory increasing with sediment depth, and reliance on water column food diminishing in tandem. (Queirós et al., 2019).
- 250 gadoids with seasonal minima both in December and March-April. The prokaryote diversity in the water column, by contrast, is lowest in the summer months. In the benthos there is no distinguishable seasonal signature to prokaryote diversity.







Figure 3: Seasonal patterns at station L4. Monthly mean values calculated across all years of available data, presented for surface (0 m) unless stated. For explanations of all data fields see Table A1. Biomass of plankton by flow cytometry derived from Table 2. Fish families plotted here are for the 11 top ranking groups based on annual mean biomass with remaining groups (including Cephalopoda) summed here as "others".



5.2 Overall seasonality: E1 and L5

Fig. 4 summarises the data available for the E1 and L5, sites further offshore than L4. All measurements except those from
the Young Fish Trawl and the shark catch data pertain to the E1 site. The Young Fish Trawl data are from site E1 and L5 combined. The shark data are from angling vessels from Looe and within 10 miles of E1.

The E1 site is more strongly stratified than L4, as evidenced by slightly higher surface temperatures and a bigger summer temperature difference between surface and depth. Being further offshore than L4 and receiving less riverine nutrient input from the rivers Tamar and Plym macronutrient concentrations are more severely limiting in summer and indeed, iron stress

- 290 from the rivers Tamar and Plym macronutrient concentrations are more severely limiting in summer and indeed, iron stress in some seasons has been suggested (Schmidt et al., 2020). This is also reflected in the stronger reduction in DIC during the stratified period, resulting in an average seasonal amplitude of 83 µmol kg⁻¹ at E1 compared to around 55 µmol kg⁻¹ at L4. Total alkalinity (TA) in contrast shows little seasonal cycle at E1 (average seasonal amplitude = 29 µmol kg⁻¹), compared to a slight increase in spring at L4 (average seasonal amplitude = 40 µmol kg⁻¹). Both L4 and E1 show a seasonal pattern of
- 295 seawater CO₂ undersaturation between January and August, followed by supersaturation in September and October, returning to near equilibrium with the atmosphere for the remainder of the year (Kitidis et al., 2012). The subsurface chlorophyll *a* maxima has been shown to be important for controlling carbon fluxes at these sites, as well as the mixing of freshwater, which is evidence by the difference between L4 and E1 conditions (Kitidis et al., 2012). The flow cytometry data reflect this (**Fig. 4**) with increased contribution of coccolithophores compared to L4, albeit with pronounced inter-annual variability and large blooms in some years but not others.
- 300 variability and large blooms in some years but not others.

Although phytoplankton and zooplankton samples are currently collected at E1 we have not summarised them in this paper because the available time series data does not cover as long a period as L4. However, a summary of phyto- and zooplankton seasonality at E1 is presented and compared with that of L4 by Djeghri (2018). These authors showed that mesozooplankton biomass at E1 is lower than at L4 and at deeper, offshore sites in the Celtic Sea (Giering et al., 2019). Also in the context of these Celtic sea stations, Schmidt (2020) examined nutrient dynamics at E1 in relation to pico- and nanoplankton, and found that late season dominance of the picocyanobacterium Synechococcus (including intense blooms in some years) tended to follow summers of particularly severe nitrate stress.

- 310 Data compiled here from the 1m², 700 µm mesh Young Fish Trawl (Fig. 4d) shows strong summer and autumn increases in pilchard eggs. The later autumn spawning period has become more dominant in recent years (Coombs et al., 2010). Fish eggs of other species, by contrast, are more abundant in early spring. *Calanus* spp. currently and historically comprise mainly *C. helgolandicus* at this site (Lindeque et al., 2013; Maud et al., 2015) and they increase in mid-summer. These large copepods are biomass-dominant mesozooplankton at the WCO (Maud et al., 2018) and important food for pelagic fish.
- 315 Success in catching blue sharks *Prionace glauca* increases rapidly until late summer.







Figure 4: Seasonal patterns at station E1+L5 (1903-2021): Monthly mean values calculated across all years of available data. Illustration shows surface (0 m) data unless stated. For explanations of all data fields see Table A2. Biomass of plankton by flow cytometry derived from Table 2.



5.3 Overall annual time trends: L4

The regular, weekly-resolved measurements at L4 are during the most recent era of rapid warming (**Fig. 2**) and both the sampling intensity and the number of planktonic taxa measured allows observation of systematic change and its driving factors. We cannot review all the literature on change here, but instead **Fig. 5** illustrates some of the key trends.

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Because stratification is such a major factor driving seasonality at the WCO, **Fig. 5** compares trends separately between the most stratified quarter of the year with lowest average nutrients (May-August) and the rest of the year. The temperature rise during the warm stratified period over the last 30 years is more pronounced than in the other months, and is well over 1°C. The sharp rise in temperature at this time of year coincides with a major decline in nitrate concentrations and DIC, pointing to the effects of enhanced stratification retarding nutrient and carbon supply (**Fig. 5a**).

Fig. 5b compares the trends for surface Chlorophyll *a* concentrations and the biomass-dominant functional groups of phytoplankton, that together dominate estimated biomass of cells counted in Lugol's preserved water samples from 10m depth, namely diatoms, dinoflagellates and nanoflagellates (ca. 15μm). Flagellates (ca. 2-15μm) are also counted more quantitatively and with full water column resolution by flow cytometry (since 2007) and the component fluorescing and non-fluorescing groups (termed nanoflagellates and heterotrophic nanoflagellates) are major contributors to community dynamics (Atkinson et al., 2021; Tarran & Bruun, 2015). The larger phytoplankton, namely diatoms, dinoflagellates and the subset of larger nanoflagellates decline strongly during the summer stratified period, with Chlorophyll *a* concentration declining overall by about 50% over 30 years.

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In parallel with these summer declines in phytoplankton, there have been declines at the same time of year in the crustacean mesozooplankton functional groups, namely large and small copepods, other crustacean holoplankton (dominated by *Podon* and *Evadne*) and fish larvae (**Fig. 5c**). The large copepod category (defined here as adult total body length over 2 mm) is strongly dominated by *Calanus helgolandicus*. From 1988 to around 2015 annual abundances of *C. helgolandicus* were fairly stable, only oscillating about four-fold between years (Atkinson et al., 2015; Maud et al., 2015). However, in recent years numbers in summer have started to decline substantially, making it hard to obtain sufficient individuals for egg production experiments. This sudden shift supports the concept of abrupt step changes that "reorganise" assemblages both at

375 Outside of summer, these declines in the crustacean groups were not seen, or were not so prevalent in other months, and there was only a weak phenological shift observed at L4 (Atkinson et al., 2015; Uriarte et al., 2021) which does not explain the differential trends between the summer and the rest of the year.

this site (Reygondeau et al., 2015) and more widely (Bode, 2023).



Other taxa, by contrast, have tended to increase at L4. Only a minority of major crustaceans have shown signs of an increase, notably the more carnivorous, late summer copepod *Centropages typicus* (Corona et al., 2021). The main increases are among meroplankton taxa, fine mesh filter feeders such as appendicularians (which dominate the "other non-crustacean zooplankton" category) as well as the gelatinous predators (dominated by cnidarians) and semi-gelatinous predators (dominated by chaetognaths). Together, this suggests a shift in the balance of the mesozooplankton, from copepod domination towards a diversity of mero- and holoplankton that are fine particle feeders, more gelatinous or more carnivorous.

These trends seen at L4 conform to much wider-scale, long-term trends that are coherent right across the NE Atlantic and NW European shelf. They are even broadly similar to those at the Naples Bay monitoring site over a similar timescale (Mazzocchi et al., 2023). As an example, the meroplankton increase is widespread across the NW European shelf and NE Atlantic (Bedford et al., 2020; Holland et al., 2023) and overall, the trends seen at L4 in summer resemble the wider trends seen particularly to the west of the UK (Schmidt et al., 2020; Holland et al., 2023). As a cause, one recent hypothesis involves a bottom-up mechanism whereby increased summer nutrient stress favours pico-size cells and cyanobacteria such as *Synechococcus* which have low polyunsaturated fatty acid (PUFA) content and poor nutritional quality. This was suggested to cause a mismatch with the energy demands of crustacean zooplankton grazers at the warmest time of year when their metabolic rates are highest (Schmidt et al., 2020). However, this does not explain the increasing abundance of carnivores or meroplankton, and to fully understand the causes of these trends, time series such as these need to be networked with those from other sites (O'Brien et al., 2017).

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Figure 5. Example plots of L4 time series showing major changes over the last 30 years. Each data point represents an annual
mean of the monthly values, averaging the main summer stratified period (May-August: red) and the rest of the year (black); (a)
Temperature, surface Nitrate plus Nitrite and DIC; (b) Phytoplankton, including surface Chl *a* concentrations and biomass of the
dominant phytoplankton functional groups counted in Lugol's-preserved water samples from 10m depth; (c) Mesozooplankton of
the "classical food web" that have declined in summer, namely crustaceans (dominated by Copepoda) and fish larvae; (d) biomass
of key taxa that have increased, including meroplankton, other holoplankton (dominated by Appendicularia), semi-gelatinous
predators (dominated by Chaetognatha) and gelatinous predators (dominated by Cnidaria). Missing months within these time
series have been replaced by overall long term mean values for the missing month. Trend lines are illustrative only, and do not

series have been replaced by overall lon necessarily imply statistical significance.



445 5.4 Overall time trends: E1

The E1 site has the longest history of measurements at the WCO and has exemplified progressive technological advances in measuring macronutrients and primary production (Southward et al., 2005). It has also been a testing ground for theories of how climatic variability impacts on nutrients and thereby on phytoplankton, cascading up to fish. The "Russell Cycle" (Cushing, 1977) was a good example of these progressive ideas, where reduced Atlantic inflows of limiting nutrients (Kemp, 1938) were suggested to reduce primary production and shifting from a herring dominated ecosystem in the first few decades of last century to a pilchard dominated one in the mid 1930s. Some of these ideas about the mechanism of the Russell cycle have since been revised (Southward et al., 2005) but nevertheless a degree of cyclicity in temperature and nutrients is clear in **Fig. 6a**, and this is manifested in major cycles of higher trophic levels (**Fig. 6b, c**).

455 A major problem when interpreting these long time series is the attribution of cause from correlative-type analyses (Bedford et al., 2020). However, the rapid rise of pilchards after the collapse of the herring fishery may be due to a combination of overfishing and climatic factors (Southward et al., 2005). Similarly, the intensive industrialised pelagic fishing for mackerel and pilchards in the 1970s and its sudden collapse due to overfishing in the 1980s has unknown effects on the trajectories of fish illustrated in **Fig. 6c**.

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- Figure 6. Examples of long time series from stations L5 and E1 from the Western Channel Observatory. Points represent averages of monthly values from the main stratified months (May-August: red) and the rest of the year (black); (a) Temperature and nutrients; (b) Catches per "standard haul" of the Young Fish Trawl, showing four categories of fish eggs and larvae with standard haul volumes standardised to a filtration volume of 4000 m³; (c) Respective panels showing: catches of *Calanus* from the Young Fish Trawl; annual mean values for total fish caught in the Young Fish Trawl (all four categories in panel b, but screened such that records with absent data for any of the four categories were removed) and then annual means calculated based on a mean of all available monthly data; Blue shark *Prionace glauca* catch per unit effort (mean catch per trip from angling boats from Looe
- 490 all available monthly data; Blue shark *Prionace glauca* catch per unit effort (mean catch per trip from angling boats from Looe fishing within 10 nautical miles of E1). Yellow bars mark the 1980s for ease of cross referencing between plots. The 1980s marked major changes, including the onset of rapid warming, the end of a period of intense pelagic trawling off Plymouth and cessation of funding for many monitoring programs including the WCO.



5.4 Calanus egg production experiments at L4

- Although rate process measurements have periodically been made at the WCO, such as primary production (Barnes et al., 2015) and grazing (Bautista & Harris, 1992; Fileman et al., 2010), measurements of *Calanus* egg production rate have been made fairly consistently since 1992 (Fig. 7). This makes it one of the longest zooplankton production time series of its kind (Harris, 2010) and offers a valuable insight both into food quantity and quality for grazers and into the population dynamics of *Calanus helgolandicus* (Green et al., 1993; Irigoien et al., 2000). While the original weekly data are archived at BODC, the monthly values averaged here (i.e. a mean of the component weekly mean rates) provide a good seasonal and long-term
- comparison with the respective monthly average water temperature and functional groups of phytoplankton.

A series of publications have used these *Calanus* egg production data and have supplied extra supporting information. Examples include the linking of egg and female condition to nutritional quality of food during the 1994 season via the use of fatty acids (Pond et al., 1996); understanding population dynamics based on the timing of egg production and the onset of stratified conditions suggested to retard egg sinking (Irigoien & Harris, 2003) and *Calanus* population dynamics in relation to food and temperature, also including measurements of egg hatch success (Bonnet et al., 2005; Cornwell et al., 2018; Maud et al., 2015; Maud et al., 2018). Long term changes in the phenology and rates of *Calanus helgolandicus* egg production have not been studied recently but these show some interesting patterns (**Fig. 7**). Until roughly 2006 there was a clear maximum of egg production per female during the spring bloom months, but over the following 15 years this moved later into the summer and autumn months, with a general decline in maximum rates. In the last couple of years, however, there are suggestions that a pattern of high egg output in spring may be re-asserting itself.

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Figure 7. Heatmap of mean egg production rate of *Calanus helgolandicus*. Pending suitable weather conditions for L4 sampling and enough adult females to incubate, experiments are run weekly with 25 female *Calanus helgolandicus* incubated for 1 day in egg production chambers and eggs then counted. Red cells: highest egg production; blue cells: lowest egg production; Cells with crosses: no data available.

5.5 Carbonate chemistry measurements

Over a decade worth of data is now available for the carbonate chemistry at both L4 and E1, which has been used to provide evidence for a number of assessments relating to ocean acidification, including the OSPAR QSR2023 OA Assessment (McGovern et al., 2023) and the recent Marine Climatic Change Impacts Partnership (MCCIP) 2022 Status on Ocean Acidification around the UK and Ireland (Findlay et al., 2022). The data series are one of just two time-series stations that record carbonate chemistry parameters in the UK at this frequency, and are submitted as part of the UKs contribution to the UN Sustainable Development Goal 14.3.1 Indicator for ocean acidification.

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Over the full time-series for L4 there has been an overall decline in both total alkalinity (TA) and dissolved inorganic carbon (DIC), which has resulted in an increase in CO_2 fugacity (fCO₂) of 6.4 µatm yr⁻¹ and a decrease in pH of -0.0126 ± 0.0022 yr⁻¹. If the 2021 data is excluded, the decrease in pH is slightly slower at -0.006 yr⁻¹ (Findlay et al. 2022), demonstrating a significant lowering of pH in 2021, a result of a decrease in alkalinity and a large reduction in salinity

545 (Gonzalez-Pola et al., 2022 monthly analysis from same sampling points as carbonate chemistry gives a decline over the

higher variability as a result of organic alkalinity contributions.



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time-series of -0.01 ± 0.005 psu y⁻¹). This rate of pH decline is faster than rates observed in the open ocean, but is similar to rates found in other near-shore locations off the French coast in the Western English Channel and Bay of Biscay (McGovern et al., 2023). Interestingly, aragonite saturation state shows no significant trend at L4, most likely resulting from the concomitant decline in both DIC and TA, but also the high level of variability in TA caused by organic alkalinity inputs from local rivers. At station E1, there has been a greater decline in DIC ($-1.35 \pm 1.12 \mu$ mol kg yr⁻¹ at E1 vs. $-0.78 \pm 0.72 \mu$ mol kg yr⁻¹ at L4), similar decline in TA and a slightly slower decline in pH (when including 2021 data: $-0.008 \pm 0.0022 \text{ yr}^{-1}$).

Since autumn 2017 additional water column measurements have been taken at L4, which provides a profile view of the carbon dynamics. As a case study, we show here the profiles between 2018 and 2020, inclusive (**Fig. 8**). There is a clear relationship between in situ density anomaly (σ t) and both DIC (DIC = 38.79. σ t + 1086, r = 0.7184, n=144, p < 0.0001) and AT (AT = 18.36. σ t +1814, r = 0.4118, n = 144, p < 0.0001). The in situ density anomaly at L4 is primarily driven by temperature, although salinity is important for the dilution of carbonate parameters at this site. The σ t represents the seasonal cycle of winter mixing followed by stratification through the spring and summer and breakdown of stratification again in the autumn. Both DIC and AT are generally at similar concentrations throughout the water column, with DIC being reduced in the upper mixed layer during stratification and corresponding to the sub-surface chlorophyll blooms (**Fig. 8**). AT has much

Data on suspended matter and particulate carbon compounds have also been collected at the WCO during different times over the years. As shown in (Rühl et al., 2021), the concentration of particulate organic carbon (POC) at a depth of 10 m at L4, measured between 2013 and 2017, is highly seasonally variable, but overall decreasing over the four-year period. It is unclear whether this is part of a more long-term cyclical pattern, or a true temporal trend in the data. Variability in particulate suspended matter concentration in general is less seasonal, and does not conform to any clear trend throughout the same time period (Rühl et al., 2021).

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Figure 8: Depth profiles through time between 2018 and 2021 of a) Chlorophyll, b) density, c) dissolved inorganic carbon (DIC), and d) total alkalinity (TA).



5.6 Benthic-pelagic coupling

- 585 Despite a clear stratification cycle, benthic dynamics of the seafloor community at L4 remain tightly linked to the water column throughout the year, as illustrated particularly in the triggering of biomass of diversity peaks and troughs linked to availability of food sources from surface waters and elsewhere (Queirós et al., 2019; Tait et al., 2015; Talbot et al., 2019). This seasonality is also observed in the dynamics of ecosystem processes mediated by macrofauna in L4 sediments i.e. bioturbation and bioirrigation (Kristensen et al., 2012) which have strong mediation effects on the rates of biogeochemical
- 590 processes at the sediment water interface, such as community respiration, and net carbon sequestration (Queirós et al., 2015; Queirós et al., 2019). Broader analyses of the seafloor time-series at L4 have also demonstrated that these dynamics are highly variable on an interannual basis, with the effects of extreme events being particularly important (Rühl et al., 2021). Net vertical flux directions of suspended and particulate matter vary throughout the year, switching in direction and respective importance for the overall flow of matter throughout the system (Rühl et al., 2020). The L4 time-series thus
- 595 provides unique value in enabling a deep assessment of trends in benthic-pelagic coupling within and across years data seldom available elsewhere is much needed to enable a deeper understanding of seafloor processes involved in biogeochemical cycles. This is especially relevant at a time when understanding, managing and enhancing the ability of the ocean to serve as a carbon sink is especially important, as it is here (at the seafloor) that long-term particulate organic carbon sequestration will take place (Williamson & Gattuso, 2022). Inter-and Intra-annual data set collections focussing on benthic-
- 600 pelagic coupling in the area have previously been published in the context of a doctoral research project (Rühl et al., 2020 and 2021).

6 Wider context

6.1 Modelling

Long-term time series like the ones reported here have been paramount in shaping our understanding of biogeochemical cycling and plankton dynamics (Benway et al., 2019). Not only have they provided the necessary data consistency to generate hypotheses to progress our understanding of marine ecosystems, they have also been critical to the advancement of our capacity to model the complex interactions between environmental and plankton dynamics. The breadth of ecosystem components that are measured routinely at the WCO has enabled a form of digital hypothesis testing using biogeochemical and plankton models (Polimene et al., 2014) comparable to the more traditional approach to hypothesis testing through experimental work under controlled laboratory conditions. The WCO timeseries have contributed to a broad range of developments of European Regional Seas Ecosystem Model (ERSEM) originating from testable hypotheses. These range from photophysiology control of plankton succession (Atkinson et al., 2018), the role of food quality on plankton blooms (Polimene et al., 2015), bacteria carbon pump (Polimene et al., 2017) or the role of mixoplankton in plankton succession dynamics (Leles et al., 2021). Models can also represent a key source of information (a concept generally referred to as data



615 augmentation) for the interpretation of time series. For example, operational models such as the Western Channel Observatory Operational Forecast model (WCOOF) (Torres & Uncles, 2011), can be used to reconstruct back trajectories of plankton samples to explain community variations or assist in the evaluation of carbon sequestration estimates (Queirós et al., 2023). Models like WCOOF can also be used to interpolate environmental conditions to explain observed plankton shifts (e.g. rapid changes to weak stratification not captured by the time-series sampling frequency) or to interpolate sparse 620 measurements (Sims et al., 2022). Ultimately, models can also inform and optimize observational approaches e.g. primary

6.2 The WCO contribution to wider observing networks to report on ocean health.

production estimation from Oxygen/Argon ratios and oxygen isotopes.

Marine time series such as those provided by the WCO form an important component to a series of wider networks for reporting on pelagic and benthic ecosystem status, and these networks span a range of scales. At the smallest scale of the SW
UK, the Western Channel Observatory observations form important contributions to the annual Southwest Marine Ecosystems Annual Reports (Smyth 2022, Atkinson et al., 2022). At the UK scale WCO data inform on regional-scale trends in plankton on the Marine Climate Change Impacts Partnership (Edwards et al., 2020; Findlay et al., 2022) and the plankton data contribute to indicator C5 within the UK's 25 Year Environment Plan. At a slightly wider scale (NW European shelf and NE Atlantic) the WCO data form part of the policy reporting to meet statutory UK policy obligations under the UK
Marine Strategy and OSPAR, for example in relation to carbonate chemistry (McGovern et al., 2023), or pelagic habitats (Ostle et al., 2021; Holland et al., 2023).

One advantage of the WCO plankton data is that they are both relatively complete in terms of taxonomic resolution and that they span multiple decades, which has enabled their use as a testbed dataset for developing indicators (McQuatters-Gollop et

- 635 al., 2019), to examine how representative single sampling stations are of wider areas (Ostle et al., 2017) and to develop indicators that include the full suite of plankton, including major groups such as gelatinous species and picoplankton, which are not included as indicators from other longer term monitoring programmes such as the Continuous Plankton Recorder. At wider ocean basin scales the Western Channel Observatory data contribute to a series of reporting networks, for example oceanography through ICES reports on ocean climate (IROC) (Gonzalez-Pola et al., 2022) or plankton through the
- 640 International Group of Marine Ecological Time series IGMETS (O'Brien et al., 2017). Because the WCO spans a small area, building these wider networks of time series is a vital tool to understand the spatial-temporal imprint of climate change amid other, more acute and localised stressors (Ratnarajah et al., 2023).

6.3 The future: melding new technological developments with existing long time series

Most of the longer time series data that we provide here have been collected with traditional techniques that require direct collection of samples, their transport to the laboratory followed by expert chemical or taxonomic analysis. These methods are expensive and time consuming and for this reason, time series worldwide are under threat from funding cuts and loss of



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expert taxonomists (Vucetich et al., 2020). Concurrent with this, new techniques are being developed for time series, for example, underway autonomous vehicles or remote collection of data with acoustic methods or moorings data are processed by automated particle imaging using machine learning particle classification, or the use of eDNA and bioinformatic processing of the enormous volumes of data collected.

Some of these newer, "big data" approaches are now being developed at the WCO. The NERC-funded Automated in-situ plankton imaging and classification system (APICS) project represents the first in-situ co-deployment of an Imaging FlowCytobot (McLane Research Laboratories, Inc.) and a Plankton Imager (Pi-10; Plankton Analytics Ltd.) in the world.
APICS will generate abundance and diversity data for organisms spanning 3-4 orders of magnitude in size, i.e., 5µm - 20mm, on hourly time-scales, which will allow a ca. 100-fold increase in phyto- and zooplankton sampling frequency at Station L4 in 2024. APICS will allow critical plankton data to be collected at the same temporal resolution as physical and chemical variables. The establishment of an in-situ imaging time-series of plankton and particles at station L4 will also facilitate the collection of highly temporally resolved pelagic SPM / POC concentration data, using image-based POC
estimation methods that are currently being refined (Giering et al., 2020) (Rühl et al. unpublished data). The WCO's rich background of contextual data, relative ease of accessibility yet at the same time its exposure to large wave amplitudes from the SW weather systems, also make it an ideal testing ground for new technology and this development is currently highly

A common conception of funders and policy makers is that new moored and autonomous instrumentation will provide a substitute for traditional monitoring that involves the collection of samples and analysis by skilled humans in a land-based laboratory. This may seem an attractive way of reducing a whole suite of costs including those for staff time, training of taxonomic skills, ship time and fuel, as well as the carbon footprint. These new methods, however, produce fundamentally different types of data to traditional approaches. This presents difficulties when melding the data together. This is a key

active, with Smartsound Plymouth (https://www.smartsoundplymouth.co.uk) providing ambitious new directions in this area.

- 670 detail, because the detection of climate change responses usually requires multiple decades of data collected in consistent fashion to have sufficient statistical power to detect change. Instead, the new approaches provide novel insights, often at much higher temporal and spatial resolution than traditional methods, better suited to capturing delicate organisms (Cross et al., 2015), vertical structure (Cornwell et al., 2020) or revealing the "hidden" diversity of assemblages through molecular metabarcoding (Lindeque et al., 2013; Parry et al., 2021). These are complementary, rather than alternatives to ongoing
- 675 monitoring and provide fresh views on how these ecosystems function. These new technologies provide far more data than can be processed manually and currently traditional methods are essential for ground-truthing the new data. We hope that sustained observations such as the WCO will embrace the strengths of both traditional and new approaches in the following decades.



7. Data availability and how to cite them

The full data and metadata are stored in a reputable UK repository known as DASSH (The Data Archive for Marine Species and Habitats) at the Marine Biological Association, Plymouth, <u>https://www.dassh.ac.uk</u>. The data are available via the link <u>http://doi.org/10.17031/645110fb81749</u>. (McEvoy & Atkinson, 2023). On using this particular version, we kindly request to cite both the actual data citation (McEvoy & Atkinson, 2023) and to cite this paper in Earth System Science Data. This paper gives a full description of the methods and correctly citing it gives due credit to the authors who contributed the datasets.
Citing the present paper when the data are used also allows standard literature searches to reveal data usage, and this provides valuable evidence to warrant continued funding of the WCO.

In future, we aim to produce more WCO data papers with updated, doi'd time series, corrections of any errors and extended data fields. Importantly, some of the older (pre-1988) time series datasets and metadata held by the MBA were not available for this data paper. We anticipate that later doi'd versions of the data will be able to include more complete historical data as well as their metadata.

8. Potential uses and limitations of these data

The dataset we provide here has both strengths and limitations. Its main strength is that it combines, for the first time, data that span from oceanography to sharks and from microbial diversity up to benthic macrofauna and fish. We hope that this is particularly valuable for education purposes: for example student projects, where the student will spend less time trying to hunt down scattered datasets and melding them together, and more time analysing them. We have also, where we could, presented the data in units of mass as well as abundance since this is particularly amenable to carbon budgets (Queirós et al., 2019), biogeochemical studies (Barnes et al., 2015) or models (Kenitz et al., 2017; Polimene et al., 2014). This study also represents the first attempt to put benthic and pelagic data sets together, and we hope this helps to make the WCO a natural laboratory to study benthic-pelagic coupling. Another advantage of this summary version of the dataset is that it spans over 100 years, with over three decades of high-quality data from multiple trophic levels. Because the WCO has witnessed substantial warming and broadly responded in a manner similar to the wider NW European shelf (Bedford et al., 2020; Schmidt et al., 2020) our summary dataset provides a test-bed to study the mechanisms that control seasonality and climate change response across multiple trophic levels.

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Our data summary also has a series of key limitations, the first one being its taxonomic resolution. To make our database manageable in size, we have condensed large species lists (over 400 planktonic taxa alone) into a just a few dozen functional groups. Users wishing to estimate diversity changes, or responses of individual key species, will need to source the original data sets via the points of contact listed on the WCO data catalogue https://westernchannelobservatory.org.uk/pelagic TS.php . Likewise, those studying short-term dynamics or "events", for





instance phenology shifts, bloom dynamics or extreme weather may prefer to access the individual timepoints which are typically weekly at L4. Despite this proviso and recommendations that <20 day resolution are needed to reveal phenology shifts (Henson et al., 2018), long-term studies of such phenomena have tended to take the pragmatic approach by averaging irregularly-spaced timepoints into monthly blocks to improve precision, data coverage and to fill data gaps (Atkinson et al., 2021; Barton et al., 2020; Edwards & Richardson, 2004; Fanjul et al., 2018; Uriarte et al., 2021).

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A final three requests to users are first: to let us know if you find any errors; second: suggest any improvements for the next version; third, please let us know if you want to incorporate this data set into a wider data networking or databasing initiative. This is to ensure that data do not become separated from metadata and that old, outdated legacy versions of the data do not linger on data portals.

8 Author contribution

AJMc and AA co-ordinated the project. AJMc compiled the monthly database from the component datasets. AA, RA, RB, IB, EF, HF, CM, AJMc, CO, PS, TS, GT, KT, ST and CW submitted the individual datasets. AJMc and AA prepared the manuscript with contributions from HF, GT, CW, SR, RT, AQ, RA, RB, EF, TS, KT, CO. All authors are either current or previous producers of the component data sets.

9 Acknowledgements

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10 Competing interests

735 The authors have no competing interests





11 Appendices

Table A1: L4 metadata for WCO monthly time-series 1988-2022. Includes sampling and analysis protocols pus data coverage and links for availability of full data sets. Column numbers reference the data sheet available via doi: 10.17031/645110fb81749

Data type	Sampling and Analysis method	Data coverage
Column headings		Availability of full data
L4: Water temperature		
L4_Temp_0m_degC L4_Temp_10m_degC L4_Temp_25m_degC L4_Temp_50m_degC L4_Temp_50m_degC	Taken weekly where conditions allow. March 1988 to April 1993 surface temperature (0m) measured using a mercury thermometer in a stainless-steel bucket of freshly collected seawater. May 1993 to Dec 2001 a PML CTD was also used	March 1988 to Dec 2021 monthly data derived from 1491 sampling points. May 1993 onwards CTD was used providing 1142, 1140 and 977 weekly
Columns 3-6	concurrently with the bucket method. Jan 2002 this PML CTD was replaced by a SeaBird SBE19+ CTD. Bucket temperatures were adjusted to CTD equivalents using a regression equation for parallel determinations. For surface values, we obtained a value for each sampling week based on this adjusted bucket temperature if only this was available. If both bucket and CTD data were available, we used the CTD temperature. We then derived arithmetic mean temperatures for each month.	timepoints for 10m, 25m and 50m respectively. https://www.westernchannelobservatory. .org.uk/data.php
L4: Nutrients		
L4. Nitrite_0m_µM L4. Nitrite_25m_µM L4. Nitrite_55m_µM L4. Nitrite_50m_µM L4. Nitrite+Nitrate_0m_µM L4. Nitrite+Nitrate_25m_µM L4. Nitrite+Nitrate_0m_µM L4. Nitrite+Nitrate_0m_µM L4. Nitrite+Nitrate_0m_µM L4. Nitrite+Nitrate_0m_µM L4. Nitrite+Nitrate_0m_µM L4. Ammonia_0m_µM L4. Ammonia_10m_µM L4. Ammonia_10m_µM L4. Ammonia_0m_µM L4. Silicate_0m_µM L4. Silicate_0m_µM L4. Silicate_10m_µM L4. Splicate_0m_µM L4. Phosphate_0m_µM L4. Phosphate_0m_µM <t< td=""><td>Taken weekly where conditions allow. Samples returned in the cool and dark to the laboratory in Plymouth as soon as possible. Triplicate samples are analysed using 0.2µm Millipore Fluoropore filtered and non-filtered water. Analyser is a 5-channel Bran+Luebbe segmented flow system. Methodology standardised according to PML protocols. Since 2007 samples analysed as soon as possible after collection. Prior to this samples were frozen and analysed in batches. Due to storage method concentrations of ammonia should be treated with care. More appropriate to consider trends rather than accurate concentrations. Quality control procedures carried out using KANSO certified reference material. Scientists participate in QUASIMEME programme. This summary data set provides a mean value of all available determinations within any given calendar month. In the original data set the symbol "<" refers to concentrations below detection limit. These have been assigned a value of zero before averaging</td><td>Surface (0m) Jan 2000 to Dec 2021 Profile (10m 25m 50m) Jan 2012 to Dec 2021 Full data lists individual replicate measurements from the weekly resolution sampling. Publicly-accessible nutrient data accessed on 14 Jul 2022 from https://www.westernchannelobservatory. .org.uk/data.php</td></t<>	Taken weekly where conditions allow. Samples returned in the cool and dark to the laboratory in Plymouth as soon as possible. Triplicate samples are analysed using 0.2µm Millipore Fluoropore filtered and non-filtered water. Analyser is a 5-channel Bran+Luebbe segmented flow system. Methodology standardised according to PML protocols. Since 2007 samples analysed as soon as possible after collection. Prior to this samples were frozen and analysed in batches. Due to storage method concentrations of ammonia should be treated with care. More appropriate to consider trends rather than accurate concentrations. Quality control procedures carried out using KANSO certified reference material. Scientists participate in QUASIMEME programme. This summary data set provides a mean value of all available determinations within any given calendar month. In the original data set the symbol "<" refers to concentrations below detection limit. These have been assigned a value of zero before averaging	Surface (0m) Jan 2000 to Dec 2021 Profile (10m 25m 50m) Jan 2012 to Dec 2021 Full data lists individual replicate measurements from the weekly resolution sampling. Publicly-accessible nutrient data accessed on 14 Jul 2022 from https://www.westernchannelobservatory. .org.uk/data.php
LA: Carbonate chemistry DIC (dissolved inorganic carbon) and TA (total alkalinity) L4_DIC_0m_µmol kg-1 L4_DIC_10m_µmol kg-1 L4_DIC_25m_µmol kg-1 L4_TA_0m_µmol kg-1 L4_TA_0m_µmol kg-1 L4_TA_25m_µmol kg-1 L4_TA_50m_µmol kg-1 Columns 27-34	 Taken weekly where conditions allow. Borosilicate glass bottles with ground glass stoppers were used to collect seawater from the Niskin bottles. Sample bottles were rinsed, filled and poisoned with mercuric chloride according to standard procedures detailed in Dickson et al. (2007). Samples were returned to PML for analysis. DIC was measured using a Dissolved Inorganic Carbon Analyser (Apollo SciTech, Model AS-C3). The analyser adds a strong acid (10% H3PO4 plus 10% NaCl solution) causing carbon species within the seawater to be converted to CO2 gas, which is purged from the sample by pure nitrogen (N2) carrier gas, is dried and cooled to reduce water vapour. The concentration of the dried CO2 gas is measured with a LICOR L1-7000 CO2 analyser. The total amount of CO2 is quantified as the integrated area under the concentration-time curve, and converted to DIC using a standard curve created by analysing known concentrations of the Certified Reference Materials (Dickson CO2 CRMs). A measurement wolume of 0.75 mL was used, with up to 5 measure standar from each sample. Values outside a 0.1 % range were excluded from the final result. Duplicate measurements provided an estimate of measurement entor < 0.1 %. DIC was corrected for the addition of mercuric chloride. TA was measured using the open-cell potentiometric titration 	Surface, 0m and 50m Oct 2008 to Dec 2020 10m and 25m Sep 2017 to Dec 2020 Data are available from British Oceanographic Data Centre (BODC) and are citable via doi:10.5285/1ec0cae5-071d-16e1-e053- 6c86abc07d47/ https://www.westernchannelobservatory .org.uk/C_chem.php



method (Dickson et al. 2007) on 12 mL sample volumes using an automated titrator (Apollo SciTech Alkalinity Titrator Model AS-ALK2). Calibration was made using Certified Reference Materials (Dickson CO2 CRMs). Duplicate measurements were made for each sample, and the estimate of measurement error < 0.5 %. TA was corrected for the addition of mercuric chloride.	
Borosilicate glass bottles with ground glass stoppers were used to collect seawater from the Niskin bottles for the methane and nitrous oxide, bott gasses were determined from the same bottle. Prior to all depths being collected in 2019 samples were collected in triplicate. Sample bottles were rinsed, filled and poisoned with mercuric chloride according to standard procedures detailed in Dickson et al. (2007). Samples were returned to PML for analysis. All samples were analysed within 3 months of collection Samples were placed into a water bath at 25°C and temperature equilibrated for a minimum of one hour before analysis.	Surface N2O coverage is from 2011 and CH4 from 2013 All 4 depths were sampled from 2019. https://www.westernchannelobservatory .org.uk/data.php
Samples were analysed by single-phase equilibration gas chromatography using a Flame Ionisation Detector for CH4, and electron capture detector for N2O and similar to that described by (Upstill-Goddard 1996). Samples were calibrated against three certified (±5%) reference standards (Air Products Ltd) which are traceable to NOAA WMO-N2O-X2006A. Concentrations in seawater at equilibration temperature (-25°C) and salinity were calculated from solubility tables of Weiss and Price(1980).	
Taken weekly where conditions allow. On each sampling date, 5L of seawater was collected from the	Feb 2012 to Nov 2019. Data are available from PML Karen
Sterives cartridge (Millipore). This was then stored at -80°C at PML before further processing. Nucleic acids were extracted using the Qiagen AllPrep DNA/RNA Mini Kit. The sterivex barrel was first filled with RLT lysis buffer and heated to 65°C for 30 mins. DNA and RNA was then extracted from the lysate following the manufacturer's instructions. DNA samples were used for microbiome analyses by sequencing of 16S rRNA genes using the Earthmicrobiome V4 PCR primers 515F (GTGYCAGCMGCCGGGGTAA) and 806R (GGACTACNVGGGTWTCTAAT). Sequencing was performed on the Miseq Personal Sequencer (Illumina, San Diego, CA, USA) using the V2 500 reagent kit by commercial contract (NU_OMICS, UK). Demultiplexed paired end FASTQ files were analysed using QIME2 and amplicon sequence variants (ASVs) generated using DADA2. For each sample, the number of ASVs (S), Pielou evenness and Shannon diversity were calculated.	Tait. https://www.westernchannelobservatory .org.uk/data.php
Taken weekly where conditions allow Triplicate 100 ml water samples filtered onto 25 mm GFF filters. Extracted overnight at 4 deg C and analysed on a Turner Fluorometer according to Welshmeyer 1994.	Surface (0m) and 10m Feb 1992 to 2020 with 1110 and 568 weekly resolution samples respectively. All depths sampled from 2018 Publicly-accessible nutrient data accessed from https://www.westernchannelobservatory
	https://www.westernchannelobservatory .org.uk/data.php
Taken weekly where conditions allow Parameter Names (if shortened versions used in column titles): [TChl a] = Total chlorophyll a = [Chlide a] + [DVChl a] + [Chl] = Total chlorophyll = [TChl a] + [TChl b] + [TChl c]; [PPC] = Photoprotective carotenoids = [Allo]+[Diad]+[Diato]+[Zea]+[Caro]; [PSC] = Photosynthetic carotenoids = [But]+[Fuco]+[Hex fuco]+Perid]; [PSP] = Photosynthetic pigments = [PSC]+[TChl]; [TAcc] = Total accessory pigments = [PPC]+[PSC]+[TChl b]+[TChl c]; [TPig] = Total pigments = [TAcc]+[TChl a].	Surface coverage is from March 1999 to Dec 2014 10, 25 and 50 m from 2009 onwards (some gaps in data) until 2014 Source data accessed via https://www.westernchannelobservatory .org.uk/data.php
	an automated tirtator (Apollo SciTech Alkalinity Tirator Model AS-ALK2). Calibration was made using Certified Reference Materials (Dickson CO2 CRMs). Duplicate measurements were made for each sample, and the estimate of measurement error < 0.5 %. TA was corrected for the addition of mercuric chloride. Borosilicate glass bottles with ground glass stoppers were used to collect seawster from the Niskin bottles for the methane and nitrous oxide, both gasses were determined from the same bottle. Prior to all depths being collected in 2019 samples were collected in triplicate. Sample bottles were rinsed, filled and poisoned with mercuric chloride according to standard procedures detailed in Dickson et al. (2007). Samples were returned to PML for analysis. All samples were analysed within 3 months of collection Samples were analysed by single-phase equilibration gas chromatography using a Flame Ionisation Detector for CH4, and electron capture detector for N2O and similar to that described by (Upstill-Goddard 1996). Samples were calibrated against three certified (±5%) reference standards (Air Products Ltd) which are traceable to NOAA WMO-N2O-X2006A. Concentrations in seawater at equilibration temperature (-25°C) and salinity were calculated from solubility tables of Weiss and Price(1980). Taken weekly where conditions allow. On each sampling date, 5L of seawater was collected from the surface and filtered immediately (on board) through a 0.22mm Sterivex cartidge (Millipore). This was then stored at -80°C at PML before further processing. Nucleic acids were extracted using the Qiagen AllPrep DNA/RNA Mini Kit. The sterivex barrel was first filled with RLT lysis buffer and heated to 55°C for 30 mins. DNA and RNA was then extracted from the lysate following the manufacturer's instructions. DNA samples were used for microbiome analyses by sequencing of Sr RNA genes using the Earthmicrobiome V4 PCR primers 515F (GTGYCAGCMCCCGCGGTAAT) and 806R (GGACTACNVGGGTWTCTAAT). Sequencing was performed on the Miseq Personal Sequencer





L4_[TPig]_10m_HPLC_mgm-3	not quantified.	
	Total chlorophyll b = chlorophyll b + divinyl chlorophyll b.	
L4_[TChl a]_25m_HPLC_mgm-3	Divinyl chlorophyll b coelutes with chlorophyll b under HPLC	
L4_[TChl]_25m_HPLC_mgm-3	conditions used to generate these data, so were not quantified	
L4_[PPC]_25m_HPLC_mgm-3	separately. Divinyl chlorophyll b is not expected to be present	
L4_[PSC]_25m_HPLC_mgm-3	in UK waters.	
L4 [PSP] 25m HPLC mgm-3	Total chlorophyll $c = chlorophyll c1 + chlorophyll c2 +$	
	chlorophyll c3	
L4_[TAcc]_25m_HPLC_mgm-3		
L4_[TPig]_25m_HPLC_mgm-3	Carotenes = $\beta \varepsilon$ -Carotene + $\beta \beta$ -Carotene	
	Alloxanthin: quantified by both HPLC methods used to	
L4_[TChl a]_50m_HPLC_mgm-3	generate L4 pigment data	
L4_[TChl]_50m_HPLC_mgm-3	19'-butanoyloxyfucoxanthin: quantified by both HPLC	
L4_[PPC]_50m_HPLC_mgm-3	methods used to generate L4 pigment data	
L4_[PSC]_50m_HPLC_mgm-3	Diadinoxanthin: quantified by both HPLC methods used to	
L4_[PSP]_50m_HPLC_mgm-3	generate L4 pigment data	
L4_[TAcc]_50m_HPLC_mgm-3	Diatoxanthin: quantified by both HPLC methods used to	
L4_[TPig]_50m_HPLC_mgm-3	generate L4 pigment data	
	Fucoxanthin: quantified by both HPLC methods used to	
Columns 50-77	generate L4 pigment data	
Columns 50 77	19'-hexanoloxyfucoxanthin: quantified by both HPLC methods	
	used to generate L4 pigment data. May include prasinoxanthin	
	(when present) for data generated using Barlow HPLC method	
	(1999-2011)	
	Peridinin: quantified by both HPLC methods used to generate	
	L4 pigment data	
	Zeaxanthin: quantified by both HPLC methods used to	
	generate L4 pigment data	
	Chlorophyll a: includes allomers and epimers: quantified by	
	both HPLC methods used to generate L4 pigment data	
	Divinyl chlorophyll a: quantified by both HPLC methods used	
	to generate L4 pigment data	
	Chlorophyllide a: quantified in 2002; 2004-5 and May 2011	
	onwards	
	Chlorophyll b and divinyl chlorophyll b: Divinyl chlorophyll b	
	coelutes with chlorophyll b under HPLC conditions used to	
	generate these data, so were not quantified separately.	
	Dinvinyl chlorophyll b is not expected to be present in UK	
	waters.	
	Chlorophyll c1: Quantified separately from chlorophyll c2	
	from May 2011 onwards.	
	Chlorophyll c2: Includes chlorophyll c1 for data from 1999-	
	April 2011	
	Chlorophyll c3: quantified by both HPLC methods used to	
	generate L4 pigment data	
	βε-carotene (alpha-carotene): quantified separately from ββ-	
	carotene from May 2011 onwards.	
	ββ-carotene (beta-carotene): includes βε-carotene for data from	
	1999-April 2011.	
	A.	
	Barlow HPLC Method reference: Barlow RG et al. (1997)	
	Column: MOS-2 Hypersil; 100x4.6mm; 3um particle size	
	Flow rate: 1mL/min	
	Mobile phase: Barlow et al. 1997	
	Extraction solvent and volume: 90% acetone; 2mL	
	Internal standard used?: Yes, Trans-B-Apo-8'-carotenal used	
	until 2008.	
	Disruption method and time: Sonication (probe), 35s	
	Soak time: 1 hr	
	Clarification procedure: Centrifugation	
	Injection procedure and volume: Autosampler	
	mixes sample with ammonium acetate (1 M) in 50/50 ratio by	
	volume. Injects 50 uL	
	Calibration Procedure: Single point	
	Source of standards: DHI, Denmark	
	Absorption coefficients used: Those provided with standards	
	by DHI	
	Expected capability of method: Not recorded	
	Quality assurance protocols: Up to 20 samples were analysed	
	per day, so maximum time of samples in autosampler is 24 h.	
	Autosampler is maintained at 4oC.	
	Zapata HPLC Method reference: Zapata M et al.	
	(2000)	
	Column: Waters C8 Symmetry; 150x2.1 mm; 3.5um particle	
	size	
	Flow rate: 200uL/min	
	Mobile phase: As described by Zapata et al. 2000	
	Extraction solvent and volume: 90% acetone; 2mL	
	Internal standard used?No	
	Disruption method and time: Sonication (probe), 35s	
	Soak time: 1 hr	
	Clarification procedure: Centrifugation and filtration (0.45 um	
	Teflon syringe filter)	
	Injection procedure and volume: Autosampler	





	mixes 200 uL sample and 80 uL water in a vial. 25 uL of this mixture is injected (actual injection volume of sample = 17.86 uL Calibration Procedure:Multipoint; three solutions bracketing the LOQ, and three bracketing the expected sample concentration Source of standards: DHI, Denmark Absorption coefficients used: Those provided with standards by DHI Expected capability of method: Average precision and accuracy for chl a (standards) was 1.44 and 2.01%, respectively Quality assurance protocols: First run of the day was discarded. A sample of mixed pigments was run prior to any samples to check retention times and resolution of critical pairs. Three samples of chlorophyll standard were analysed with each sample set to check response factor is within 5% of calibration value. Up to 20 samples are analysed per day, so maximum time of samples in autosampler is 24 h. Autosampler was maintained at 4oC. Pipette accuracy determined daily by weighing.	
14: <20 µm plankton abundance profiles measured by flow cytometry 14. Syn_0m_FCM_cells mL-1 14. Picoeuk_0m_FCM_cells mL-1 14. Nanoeuk_0m_FCM_cells mL-1 14. Crypto_0m_FCM_cells mL-1 14. Crypto_0m_FCM_cells mL-1 14. HNApacteria_0m_FCM_cells mL-1 14. HNApacteria_0m_FCM_cells mL-1 14. HNApacteria_0m_FCM_cells mL-1 14. JNApacteria_0m_FCM_cells mL-1 14. Napacteria_10m_FCM_cells mL-1 14. Napacteria_10m_FCM_cells mL-1 14. HNApacteria_10m_FCM_cells mL-1 14. HNApacteria_10m_FCM_cells mL-1 14. JNApacteria_10m_FCM_cells mL-1 14. JNApacteria_10m_FCM_cells mL-1 14. JNApacteria_10m_FCM_cells mL-1 14. Syn_25m_FCM_cells mL-1 14. Syn_25m_FCM_cells mL-1 14. Syn_25m_FCM_cells mL-1 14. JNApacteria_25m_FCM_cells mL-1 14. JNApacteria_50m_FCM_cells mL-1 <td>Taken weekly where conditions allow Analysed in triplicate (phytoplankton and bacteria) or duplicate (heterotrophic nanoflagellates). Vertical profiles of the mean abundance of groups of microbial plankton as cells per millitilire, measured using flow cytometry (BD Accuri C6 flow cytometer) The groups quantified are divided into phytoplankton and heterotrophs. Phytoplankton groups quantified are: Syn Synechococccus sp. (cyanobacteria) Picoeuk Picoeukaryotes (smaller than 3 μm) Crypto Cryptophytes Cocco Coccolithophores Nanoeuk Nanoeukaryotes not already mentioned (2-20 μm). Heterotrophs quantified are: HNan heterotrophic hanoflagellates HNAbacteria heterotrophic bacteria with relatively high nucleic acid content LNAbacteria heterotrophic bacteria with relatively low nucleic acid content.</td> <td>April 2007 to Dec 2021 Source data accessed via https://www.westernchannelobservatory .org.uk/data.php</td>	Taken weekly where conditions allow Analysed in triplicate (phytoplankton and bacteria) or duplicate (heterotrophic nanoflagellates). Vertical profiles of the mean abundance of groups of microbial plankton as cells per millitilire, measured using flow cytometry (BD Accuri C6 flow cytometer) The groups quantified are divided into phytoplankton and heterotrophs. Phytoplankton groups quantified are: Syn Synechococccus sp. (cyanobacteria) Picoeuk Picoeukaryotes (smaller than 3 μm) Crypto Cryptophytes Cocco Coccolithophores Nanoeuk Nanoeukaryotes not already mentioned (2-20 μm). Heterotrophs quantified are: HNan heterotrophic hanoflagellates HNAbacteria heterotrophic bacteria with relatively high nucleic acid content LNAbacteria heterotrophic bacteria with relatively low nucleic acid content.	April 2007 to Dec 2021 Source data accessed via https://www.westernchannelobservatory .org.uk/data.php
L4: Microscopy analysis of lugols and formalin preserved phytoplankton L4. Diatoms_10m_microscopy_cells ml-1 L4_Dinoflagellates_10m_microscopy_cells ml-1 L4_Flagellates_10m_microscopy_cells ml-1 L4_Flagellates_10m_microscopy_cells ml-1 L4_Flagellates_10m_microscopy_cells ml-1 L4_Diadoms_10m_microscopy_cells ml-1 L4_Diadoms_10m_microscopy_cells ml-1 L4_Diatoms_10m_microscopy_mgC m-3 L4_Dinoflagellates_10m_microscopy_mgC m-3 L4_Diadoms_10m_microscopy_mgC m-3 L4_Flagellates_10m_microscopy_mgC m-3 L4_Piaeocystis_10m_microscopy_mgC m-3 L4_Diadoms_10m_microscopy_mgC m-3 L4_Ciliates_10m_microscopy_mgC m-3	Taken weekly where conditions allow Paired 200mL water samples collected from 10m depth using Niskin bottle attached to the CTD are immediately fixed in 1) acid Lugol's iodine (for all taxa except coccolithophores) and 2) neutral formaldehyde for coccolithophores. Sub samples are analysed by light microscopy using the settlement technique (Utermohl, 1958) and identified to species level where possible. Organised into six functional groups. Mean cell dimensions of each taxa are used to calculate species-specific biovolumes which are converted to carbon biomass using the equations of (Menden-Deuer & Lessard, 2000)	Single depth (10m) October 1992 – December 2020, except for gaps in sampling between October 1994 – May 1995 and December 2011 Data are available from British Oceanographic Data Centre (BODC) and are citable via https://www.bodc.ac.uk/data/published_ data_library/catalogue/10.5285/c9386b5 c-b459-782f-e053-6c86abc0d129/ https://www.westernchannelobservatory .org.uk/data.php





	Abundance data are presented as cells per mL and biomass as mgC per m3. Note: In 2005 sample collection was via a deck hose. This caused damage to the fragile ciliates hence the count is much lower for that year.	
14: FlowCam analysis of 63µm mesh plankton net hauls (50-0m) 14_Total Diatoms_FlowCam_mgCm-3 14_Total Dinoflagellates_FlowCam_mgCm-3 14_Colony flagellates_FlowCam_mgCm-3 14_Large Protists_FlowCam_mgCm-3 14_Total Copepod nauplii_FlowCam_mgCm-3 Columns 122-127	Taken weekly where conditions allow Water samples collected from a 0-50m vertical haul using a 63µm mesh WP2 style net (UNESCO, 1968, p. 153–157). Mesh change in July 2019 from 63µm to 50µm. Prior to analysis samples are pre-screened using a 300µm mesh. However, net samples collected between June 2015 and May 2016 were pre-screened using a 200µm mesh. Sample analysed live whenever possible using a FlowCam VS IV model fitted with a 300µm flowcell. Analysis carried out using x4 magnification using auto-image mode. Classification of acquired images carried out using Visualspreadsheet (2012-2016) and Ecotaxa (2017-2019). Taxa were then assigned to six broad functional groups. Mean cell dimensions of each taxa were used to calculate species-specific biovolumes which were converted to carbon biomass using suitable C conversion equations. Biomass is presented as mgC per m3. For Diatoms, Dinoflagellates (excluding Noctiluca, & Neoceratium spp) and Clilates, morphological information and shape assignment was used to calculate biovolume (Alvarez et al 2012 Table 1.). For Noctiluce and Neoceratium spp., mean cell volumes were taken from Widdicombe et al (2010). For all other Dinoflagellates, Diatoms and Clilates, cell biovolumes were converted to carbon biomass using the equations of Menden- Deuer and Lessard (2000). For large protists mostly Radiolaria, the C conversion in Michaels et al (1995) was used. Colonial flagellates were converted to C according to Børsheim & Brabak, (1987). Biomass of Copped nauplii was calculated using the equations of Uye et al (1996).	Sept 2012 to Dec 2013 are from 43 time points June 2015 to Dec 2019 are from 163 time points Abundance data are also available for meroplankton taxa, these have not been converted to biomass to date. Source data accessed via https://www.www.testernchannelobservatory .org.uk/data.php
L4: Noctiluca scintillans microscopy analysis of WP2 net hauls (50-0m) L4_Noctiluca scintillans_WP2net_no.m-3 L4_Noctiluca scintillans_WP2net_mcgC.m-3 Columns 128-129	Taken weekly where conditions allow. Two vertical hauls (50-0m) are taken using 200 micron WP2 nets (UNESCO, 1968, p. 153–157) Both replicates samples are analysed by subsampling, enumerated and identified, currently using a Olympus SZX16 stereo microscope fitted with a SZX2-ILLT LED transmitted light illuminator stand. Source data represents weekly average abundance across the two replicates and converted to numbers in a m3. Monthly abundance represent an arithmetic mean value from between 1 and 5 visits in any given month and on a weekly basis. Biomass calculations derived from abundance data using a conversion factor of 0.020375mcgC per cell using the equations of (Menden-Deuer & Lessard, 2000). Be aware that zeros are present from 2009 onwards where there is confidence in the data. A zero represents looked for but not present in the sample analysed. Data prior to this is less certain so zeros have been omitted.	July 1997-2021. Source data available via McEvoy A.; Atkinson A.; Beesley A.(2022). Zooplankton abundance time series from net hauls at site L4 off Plymouth, UK between 1988-2021. https://www.bodc.ac.uk/data/published_ data_library/catalogue/10.5285/c785f2f 7-05d5-2f47-e053-6c86abc08bee/ https://www.westernchannelobservatory .org.uk/data.php



L4: Zooplankton microscopy analysis of WP2 net		
L4: Zooplankton microscopy analysis of WP2 net hauls (50-0m) L4_meroplankton_WP2net_no.m-3 L4_ismall_copepods_WP2net_no.m-3 L4_lispe_copepods_WP2net_no.m-3 L4_ispe_topepods_WP2net_no.m-3 L4_gelatinous_predators_WP2net_no.m-3 L4_semi-gelatinous_predators_WP2net_no.m-3 L4_other_non-crustacean_holoplankton_WP2net_no.m-3 L4_other_non-crustacean_holoplankton_WP2net_no.m-3 L4_meroplankton_WP2net_mgCm-3 L4_fish_larvae_WP2net_mgCm-3 L4_gelatinous_predators_WP2net_mgCm-3 L4_gelatinous_predators_WP2net_mgCm-3 L4_gelatinous_predators_WP2net_mgCm-3 L4_gelatinous_predators_WP2net_mgCm-3 L4_other_crustacean_holoplankton_WP2net_mgCm-3 L4_other_crustacean_holoplankton_WP2net_mgCm-3 L4_other_non_crustacean_holoplankton_WP2net_mgCm-3 L4_other_non_crustacean_holoplankton_WP2net_mgCm-3 L4_other_non_trustacean_holoplankton_WP2net_mgCm-3 L4_other_non_trustacean_holoplankton_WP2net_mgCm-3 L4_other_non_trustacean_holoplankton_WP2net_mgCm-3 L4_other_non_trustacean_holoplankton_WP2net_mgCm-3 L4_other_non_trustacean_holoplankton_WP2net_mgCm-3 L4_other_non_trustacean_holoplankton_WP2net_mgCm-3 L4_other_non_trustacean_holoplankton_WP2net_mgCm-3 <	Taken weekly where conditions allow. Two vertical hauls (50-0m) are taken using 200-micron WP2 nets (UNESCO, 1968) Both replicates' samples are analysed by subsampling, enumerated and identified currently using an Olympus SZX16 stereo microscope fitted with a SZX2-1LLT LED transmitted light illuminator stand. More details are provided in Atkinson et al (2015). Source data comprises average abundance of the taxa that have been consistently identified since 1988. These source data are weekly averages across the two replicates, converted to numbers per m3 and biomass estimated. Data presented here have been aggregated into functional groups broadly based on the lifeforms for policy reporting in Ostle et al (2021). There has, however, been a few further subdivisions to better reflect trophic mode. These functional group allocations are coded, and numbers-to-biomass conversion factors are provided within the "trait" header bar data from the source dataset. Therefore there are 8 functional groups based partly on size, taxonomy and trophic mode, and with separate columns for abundance and estimated biomass. Because biomass is a derived property, often with different conversion factors between the four seasons (see data source doi), it is best to use numerical abundance data for population dynamics studies and biomass data for models, carbon budgets etc. As previously stated the groups comprise the whole of the consistently identified zooplankton so adding them will give a good estimate of total metazoan zooplankton with the exception of Ctenophores (see below).	March 1988 to Dec 2020. Derived from 1452 sampling timepoints with a weekly resolution. Monthly mean data are available for each intervening month except August 2000 and typically represent an arithmetic mean value across between 1 and 5 weekly visits in any given month. McEvoy, A., Beesley, A., & Atkinson, A. (2022). Subset of zooplankton abundance and biomass time series from net hauls at site L4 off Plymouth, UK between 1988-2020. (Version 1) [Data set]. NERC EDS British Oceanographic Data Centre NOC, https://doi.org/10.5285/D7FB6C E3-7BC9-307B-E053-6C86ABC0671B https://www.westernchannelobservatory .org.uk/14_zooplankton.php
	biomass data for models, carbon budgets etc. As previously stated the groups comprise the whole of the consistently identified zooplankton so adding them will give a	
	GELATINOUS PREDATORS: Cnidarians only, dominated in terms of numbers and biomass by Siphonophores. A notable taxon not included is Ctenophores, due to potential inconsistency in counting in early years and due to preservation issues, which we are in the process of resolving. SEMI-GELATINOUS PREDATORS: Chaetognaths and <i>Tomopteris</i> spp., with numbers and biomass strongly dominated by the former. OTHER CRUSTACEAN HOLOPLANKTON: These are the remaining groups of crustacean holoplankton not covered above, namely <i>Evadue</i> spp. <i>Podon</i> spp., Hyperiidae amphipods, mysids, and the various naupli to adult stages of Euphausitids. They are strongly dominated numerically and in terms of biomass by the Cladocerans (miscoded as non- crustaceans in the source file). OTHER NON-CRUSTACEAN HOLOPLANKTON: These are the remaining groups of crustacean holoplankton not covered above, namely Appendicularians, <i>Limacina</i> spp., Doliolids and <i>Clione</i> spp. They are strongly dominated numerically and in terms of biomass by Appendicularians.	
L4: Calanus helgolandicus weekly egg production using females from the Western English Channel site L4 L4_Calanus eggs_watercolumn_expt_eggs per female per day	Taken weekly where conditions allow A live sample is collected and returned in the cool and dark to the laboratory in Plymouth as soon as possible. Sample is gently poured through a 200-micron mesh sieve <i>Calanus sp</i> females in healthy condition are picked out gently	Feb 1992 to Nov 2021 where availability of <i>Calanus</i> females allow. Data represents Mean No eggs per female per day.
Column 146	using stork-billed forceps under a microscope as quickly as possible. Five replicates each containing 5 female <i>Calanus</i> sp are	McEvoy A.; Beesley A.; Atkinson A.(2022). Calanus helgolandicus



	incubated in the dark in filtered seawater for 24 hours. Each	weekly egg production time series
	beaker contains an egg collector. Temperature follows ambient conditions at L4 surface. The eggs produced are collected and counted. Females are identified for species. Eggs retained for hatching success.	between 1992-2021, using females from the Western English Channel site L4. https://www.westernchannelobservatory .org.uk/calanus_egg_production.php
L4: Sediment 16S alpha diversity L4_sediment_prokayote_diversity_S_0m_16S SEQ L4_sediment_prokayote_diversity_Pielou_0m_16s SEQ L4_sediment_prokayote_diversity_Shannon_0m_16s SEQ Columns 147-149	Sediments are collected using a box corer and the uppermost 0 - Icm carefully sampled by scraping into a sterile 2mL tube. Eight replicate samples are taken for each sampling time, and four of these are used for DNA extraction using 0.5g sediment and Qiagen's DNeasy PowerSoil Kit according to the manufacturer's instructions. 165 rRNA genes were sequenced using the Earthmicrobiome V4 PCR primers 515F (GTGYCAGCMGCCGCGGTAA) and 806R (GGACTACNVGGGTWTCTAAT). Sequencing was performed on the MiSeq Personal Sequencer (Illumina, San Diego, CA, USA) using the V2 500 reagent kit by commercial contract (NU_OMICS, UK). Demultiplexed paired end FASTQ files were analysed using QIIME2 and amplicon sequence variants (ASVs) generated using DADA2. For each sampling occasion, the mean number of ASVs for the four replicates is calculated (S), along with Pielou evenness and Shannon diversity.	Feb 2012 to 2019. In 2012 and from 2014 to 2019, the aim was to sample monthly when possible. In 2013 samples were collected in February and September only. Data are available from PML Karen Tait. https://www.westernchannelobservatory .org.uk/data.php
 L4: Benthic fauna from box cores L4_Macrofaunal Deposit Feeders_50m_0.1m3 Box Core_Average abundance of individual taxa per month L4_Macrofaunal Suspension Feeders_50m_0.1m3 Box Core_Average abundance of individual taxa per month L4_Macrofaunal Predators_50m_0.1m3 Box Core_Average abundance of individual taxa per month L4_Macrofaunal Scavengers_50m_0.1m3 Box Core_Average abundance of individual taxa per month L4_Macrofaunal Scavengers_50m_0.1m3 Box Core_Average abundance of individual taxa per month L4_Macrofaunal Deposit Feeders_50m_0.1m3 Box Core_Average biomass of individual taxa per month L4_Macrofaunal Suspension Feeders_50m_0.1m3 Box Core_Average biomass of individual taxa per month L4_Macrofaunal Predators_50m_0.1m3 Box Core_Average biomass of individual taxa per month L4_Macrofaunal Predators_50m_0.1m3 Box Core_Average biomass of individual taxa per month L4_Macrofaunal Scavengers_50m_0.1m3 Box Core_Average biomass of individual taxa per month L4_Macrofaunal Souvengers_50m_0.1m3 Box Core_Average biomass of individual taxa per month L4_Macrofaunal Scavengers_50m_0.1m3 Box Core_Average biomass of individual taxa per month L4_Macrofaunal Souvengers_50m_0.1m3 Box 	 Taken monthly where conditions allow. 4 or 5 replicate 0.1m3 box cores of sediment collected from 50m depth. All sediment collected is sieved over a 0.5mm mesh and retained fauna preserved in 10% formaldehyde solution. Source taxa are identified and counted using stereo and compound microscopy to species level or lowest possible taxonomic resolution. Abundance and blotted wet weight (0.00000g) per taxa is recorded per 0.1m3 box core sample. From 4 principle feeding traits, based on information primarily gathered from the BIOTIC database, 1 unique principle trait was assigned per taxa; calculated using algorithms based upon body composition, maximum length and body mass. (MarLIN 2006) 	July 2008 to July 2019. Abundance and biomass data from 65 time points is presented as monthly averages per corresponding feeding trait; suspension feeders, deposit feeders, scavengers, carnivores. Mesher T., McNeill C.L. (2022). Benthic Survey Macrofauna Abundance and Biomass Data, as part of the Western Channel Observatory, UK, between 2008 and 2019. https://www.bodc.ac.uk/data/published_ data_library/catalogue/10.5285/d9f4420 2-b0d4-646c-e053-6c86abc018c6/ https://www.westernchannelobservatory .org.uk/data.php
L4: Cephalopoda and Demersal fish families by trawling L4: Cephalopoda abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Bothidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Callionymidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Callionymidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Captoidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Cepolidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Cripcidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Cippeidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Cippeidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Cippeidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Pieuronectidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Pieuronectidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Merlucciidae abundance_50-60m_Standard Haul_individuals.trawl-1 L4: Caragigidae abundance_50-60m_Standard Haul_individuals.trawl-1	Standard hauls were collected using a large otter trawl (2008- June 2014), a Channel Hunter box trawl (July 2014-March 2015) deployed from Plymouth Quest, then a modified Channel Hunter box trawl (April 2015-September 2018) deployed from MBA Sepia. Trawl duration was approximately 40 minutes. Only trawls from 50-60m are included. Individuals were identified to species, measured (mm) and weighed (g) on-board. Where a species was abundant a subsample was weighed and total biomass extrapolated. Abundances and biomass are reported at the family level, and only families comprising at least 1% contribution in at least one month are included.	April 2008 to Sept 2018. Between 1 and 7 trawls were collected per month sampled (total 282, average 2.88) Source data for 2015-2018 via https://portal.medin.org.uk/portal/start.p hp#detail%tpc=010_0370af22f970a98e 2a5fcc79d5dd05b1





L4_Zeidae abundance_50-60m_Standard	
Haul_individuals.trawl-1	
L4_Gobiidae abundance_50-60m_Standard	
Haul_individuals.trawl-1	
L4_Scombridae abundance_50-60m_Standard	
Haul_individuals.trawl-1	
L4_Scyliorhinidae abundance_50-60m_Standard	
Haul_individuals.trawl-1	
L4_Scophthalmidae abundance_50-60m_Standard	
Haul individuals.trawl-1	
L4 Cephalopoda biomass 50-60m Standard	
Haul_g.trawl-1	
L4_Bothidae biomass_50-60m_Standard	
Haul g.trawl-1	
L4_Soleidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Callionymidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Cepolidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Triglidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Clupeidae biomass_50-60m_Standard	
Haul g.trawl-1	
L4_Engraulidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Pleuronectidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Gadidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Merlucciidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Mullidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Carangidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Zeidae biomass_50-60m_Standard Haul_g.trawl-	
1	
L4_Scombridae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Scyliorhinidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Lophiidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Triakidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Scophthalmidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Rajidae biomass_50-60m_Standard Haul_g.trawl-	
1	
L4_Lotidae biomass_50-60m_Standard Haul_g.trawl-	
1	
L4_Moronidae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Congridae biomass_50-60m_Standard	
Haul_g.trawl-1	
L4_Squalidae biomass_50-60m_Standard	
Haul_g.trawl-1	
Ť	
Columns 158-200	



Table A2: E1 metadata for WCO monthly time-series 1903-2021. Includes sampling and analysis protocols pus data coverage and750links for availability of full data sets. Column numbers reference the data sheet available via doi :10.17031/645110fb81749

Data type	Sampling and Analysis method	Data coverage
Data type Column headings	Sampling and Analysis method	Data coverage Availability of full data
E1: Water temperature E1_Temp_0m_DegC E1_Temp_10m_DegC E1_Temp_20m_DegC E1_Temp_40m_DegC E1_Temp_40m_DegC E1_Temp_50m_DegC E1_Temp_60m_DegC E1_Temp_70m_DegC E1_Temp_70m_DegC Columns 3-10	For the early period of the E1 time-series reversing thermometers were used. Values are derived from Niskin bottles and CTD except for the period December 1985 to April 2002 when no in situ sampling was undertaken and satellite sea surface temperature data pertaining to the middle of each month were used instead . Where multiple sampling timepoints existed for a calendar month we used the arithmetic mean value. Post 2002 a SeaBird SBE19+ was used.	 1903 data begins 1910 to 1920 no data 1939 to 1945 no data Surface data are most extensive. For each depth, number of sampling timepoints were 1146, 954, 892, 609, 740, 908, 262, and 815 respectively. Source dataset was produced for the ICES Report on Ocean by Tim Smyth https://ocean.ices.dk/core/iroc https://www.westernchannelobservatory.org.uk/data.php
El_Nitrite_0m_µm El_Nitrite_10m_µm El_Nitrite_20m_µm El_Nitrite_40m_µm El_Nitrite_40m_µm	Taken fortnightly where conditions allow. Data from 2002: Samples returned in the cool and dark to the laboratory in Plymouth. Samples are stored for 2-3 hours before returning for analysis and sometimes frozen.	Jan 1934 a few records for Phosphate April 1948 records begin again for Phosphate Jan 1951 records begin for Silicate Jan 1966 records begin again for Nitrite+Nitrate Jan 1986 to Dec 2001 no data Jan 2002 to Oct 2021 all covered
E1_Nitrite_60m_µm E1_Nitrite+Nitrate_0m_µm	Triplicate samples are analysed using 0.2µm Millipore Fluoropore filtered and non-filtered water. Analyser is a 5-channel Bran+Luebbe segmented flow system.	Full data lists individual replicate measurements from the weekly resolution sampling
E1_Nitrite+Nitrate_10m_µm E1_Nitrite+Nitrate_20m_µm E1_Nitrite+Nitrate_30m_µm E1_Nitrite+Nitrate_40m_µm E1_Nitrite+Nitrate_60m_µm	Methodology standardised according to PML protocols. Due to storage method concentrations of Ammonia should be treated with care. More appropriate to consider trends rather than accurate concentrations. Quality control procedures carried out using KANSO certified reference material.	Publicly-accessible nutrient data accessed on 14 Jul 2022 from https://www.westernchannelobservatory.org.uk/data.php
El_Ammonia_Om_µm El_Ammonia_10m_µm El_Ammonia_20m_µm El_Ammonia_20m_µm El_Ammonia_40m_µm El_Ammonia_60m_µm El_Silicate_00m_µm El_Silicate_10m_µm El_Silicate_30m_µm El_Silicate_30m_µm El_Silicate_60m_µm El_Phosphate_10m_µm El_Phosphate_10m_µm El_Phosphate_20m_µm El_Phosphate_20m_µm El_Phosphate_40m_µm El_Phosphate_40m_µm	Scientists participate in QUASIMEME programme Data from last century: Data obtained from the link on the data page of the Western Channel Observatory website and extracted from the NOWESP (North West European Shelf Program) database for the period 1934-1987. Source data includes profile data from 0-80m. This monthly data uses depths 0, 10 and 20m because these are compatible with post 2002 records. Nitrite+Nitrate column header describes post 2002 records. It is unclear if the last century values refer strictly to Nitrate only or Nitrite+Nitrate. This summary data set provides a mean value of all available determinations within any given calendar month. In the original data set the symbol "<" refers to concentrations below detection limit. These have been assigned a value of zero before averaging.	
Columns 11-40		
E1: Carbonate chemistry DIC (dissolved inorganic carbon) and TA (total alkalinity) E1_DIC_0m_micromol kg-1 E1_DIC_60m_micromol kg-1	Taken fortnightly where conditions allow. Borosilicate glass bottles with ground glass stoppers were used to collect seawater from the Niskin bottles. Sample bottles were rinsed, filled and poisoned with mercuric chloride according to standard procedures detailed in	Surface 0m and 60 m depth coverage from October 2008 to December 2020. Source data set available via
E1_TA_0m_micromol kg-1 E1_TA_60m_micromol kg-1	Dickson et al. (2007). Samples were returned to PML for analysis.	https://www.westernchannelobservatory.org.uk/C_chem.php
Columns 41-44	DIC was measured using a Dissolved Inorganic Carbon Analyser (Apollo SciTech, Model AS-C3). The analyser adds a strong acid (10% H3PO4 plus 10% NaCl solution) causing carbon species within the seawater to be converted to CO2 gas, which is purged from the sample by pure nitrogen (N2) carrier gas, is dried and cooled to reduce water vapour. The concentration of the dried CO2 gas is measured with a LICOR LL7000 CO2 analyser. The total amount of CO2 is quantified as the integrated area under the concentration-time curve and converted to DIC using a standard curve created by analysing known concentrations of the Certified Reference Materials (Dickson CO2 CRMs). A measurement volume of 0.75 mL was used, with up to 5 measurements made	



	from each sample. Values outside a 0.1 % range were excluded from the final	
	result. Duplicate measurements provided an estimate of measurement error < 0.1 %. DIC was corrected for the addition of mercuric chloride.	
	TA was measured using the open-cell potentiometric titration method (Dickson et al. 2007) on 12 mL sample volumes using an automated titrator (Apollo SciTech Alkalinity Titrator Model AS-ALK2). Calibration was made using Certified Reference Materials (Dickson CO2 CRMs). Duplicate measurements were made for each sample, and the estimate of measurement error < 0.5 %. TA was corrected for the addition of mercuric chloride.	
E1: <20 µm plankton abundance profiles measured		
by flow cytometry E1_Syn_0m_FCM_ cells mL-1 E1_Picoeuk_0m_FCM_ cells mL-1	Taken fortnightly where conditions allow Most analysed in triplicate (phytoplankton and bacteria) for surface (0m) and single samples for all other depths.	March 2014 to Oct 2021 Source data set available via
E1_Nanoeuk_0m_FCM_cells mL-1 E1_Cocco_0m_FCM_cells mL-1	Vertical profiles of the mean abundance of groups of microbial plankton as cells per millilitre, measured using flow cytometry. (BD Accuri C6 flow cytometer) The groups quantified are divided into phytoplankton and heterotrophs.	https://www.westernchannelobservatory.org.uk/data.php
E1_Crypto_0m_FCM_cells mL-1 E1_HNAbacteria_0m_FCM_cells mL-1 E1_LNAbacteria_0m_FCM_cells mL-1	Phytoplankton groups quantified are: Syn: Synechococcus sp. (cyanobacteria) Picoeuk: Picoeukaryotes (smaller than 3 μm)	
E1_Syn_10m_FCM_cells mL-1 E1_Picoeuk_10m_FCM_cells mL-1	Crypto: Cryptophytes Cocco: Coccolithophores	
E1_Nanoeuk_10m_FCM_cells mL-1 E1_Cocco_10m_FCM_cells mL-1 E1_Crypto_10m_FCM_cells mL-1	Nanoeuk: Nanoeukaryotes not already mentioned (2-20 µm). Heterotrophs quantified are:	
E1_HNAbacteria_10m_FCM_cells mL-1 E1_LNAbacteria_10m_FCM_cells mL-1	HNAbacteria: heterotrophic bacteria with relatively high nucleic acid content LNAbacteria: heterotrophic bacteria with relatively low nucleic acid content.	
E1_Syn_20m_FCM_cells mL-1 E1_Picoeuk_20m_FCM_cells mL-1 E1_Nanoeuk_20m_FCM_cells mL-1 E1_Ccco_20m_FCM_cells mL-1 E1_Crypto_20m_FCM_cells mL-1 E1_HNAbacteria_20m_FCM_cells mL-1 E1_LNAbacteria_20m_FCM_cells mL-1		
E1_Syn_30m_FCM_cells mL-1 E1_Picoeuk_30m_FCM_cells mL-1 E1_Nanoeuk_30m_FCM_cells mL-1 E1_Cocco_30m_FCM_cells mL-1 E1_Crypto_30m_FCM_cells mL-1 E1_HNAbacteria_30m_FCM_cells mL-1 E1_LNAbacteria_30m_FCM_cells mL-1		
E1_Syn_40m_FCM_cells mL-1 E1_Picoeuk_40m_FCM_cells mL-1 E1_Nanoeuk_40m_FCM_cells mL-1 E1_Cocco_40m_FCM_cells mL-1 E1_Crypto_40m_FCM_cells mL-1 E1_HNAbacteria_40m_FCM_cells mL-1 E1_LNAbacteria_40m_FCM_cells mL-1		
E1_Syn_60m_FCM_cells mL-1 E1_Picoeuk_60m_FCM_cells mL-1 E1_Nanoeuk_60m_FCM_cells mL-1 E1_Cocco_60m_FCM_cells mL-1 E1_Crypto_60m_FCM_cells mL-1 E1_HNAbacteria_60m_FCM_cells mL-1 E1_LNAbacteria_60m_FCM_cells mL-1		
Columns 45-86		
E1+L5: combined Young Fish Trawl (YFT) E1+L5_ Calanus sp_YFT_No.4000m-3 E1+L5_Pilchard eggs_YFT_No.4000m-3 E1+L5_Other fish eggs_YFT_No.4000m-3	Although net design and methods of deployment have changed on several occasions, care has been taken to ensure that sampling characteristics have not altered appreciably. The 1m ² Young Fish Trawl (YFT) fitted with a 700μm knitted mesh is hauled for 20 min in an oblique profile to an ideal depth of~5m above the seabed.(Ostle et al., 2021)	1924–1940 1945–1987 2001–2013
E1+L5_Other Ish eggs_YF1_No.4000m-3 E1+L5_Clupeidae larvae_YFT_No.4000m-3 E1+L5_Other fish larvae_No.4000m-3	The samples are preserved in 4% buffered formalin and analysed as soon as possible after collection using a WILD M5 binocular microscope. The volume of filtered water is calculated using flow data recorded by a	Source data available https://doi.mba.ac.uk/data/1536
Columns 87-91	flowmeter fitted across the net mouth. Results are standardised to the number of individuals per 4000m3 in order to mitigate historical changes in sampling gear and deployment. A comprehensive summary of these macroplankton sampling methods and analysis is given in Southward et al. (2005)	
	Note: Please be aware of zero values within this dataset, generally these are true zeros but not necessarily for all. This is being checked and will be addressed in future versions of the dataset.	



E1: Recreational captures of blue shark (Prionace		
glauca)	The Pat Smith database is a collaboration between the Shark Angling Club of	1958- 1971
	Great Britain (SACGB) and the Sportfishing Club of the British Isles (SCBI).	1997-2021
E1_Prionace glauca captures_recreational anglers out of	It is a collation of information records kept by the SACGB.	
Looe_individuals	Recreational angling trips from the port of Looe, Cornwall, within 10nm radius	Annual data are available for all years 1953-2022 via pre-
E1_Prionace glauca catch per unit effort_recreational	of E1.	print
anglers out of Looe_captures.trip^-1	The data presented here are for years when monthly log-book information is	Simon Thomas et al(2023)
	currently available.	
Columns 92-93	The data record 64287 captures from 32906 trips from the port in 200 monthly	
	periods between 1958 and 2021.	
	Since 1998 all captures have been released.	
	Data presented are the total number of captures in a given month, and the	
	average catch per unit effort (as captures per trip).	

12 References

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