Upper ocean flux of biogenic calcite produced by the Arctic planktonic foraminifera *Neogloboquadrina pachyderma*

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Abstract. With ongoing warming and sea ice loss, the Arctic Ocean and its marginal seas could become less challenging to pelagic calcifiers, resulting in modifications of the regional carbonate cycle and the composition of the seafloor sediment. A substantial part of the pelagic carbonate production in the Arctic is due to the calcification of the dominant planktonic foraminifera species *Neogloboquadrina pachyderma*. To quantify carbonate production and loss in the upper water layer by this important Arctic calcifier, we compile and analyse data from vertical profiles in the upper water column of shell number concentration, shell sizes and weights of this species across the Arctic region during summer. Our data is inconclusive on whether the species performs ontogenetic vertical migration throughout its lifecycle, or whether individual specimens calcify at a fixed depth within the vertical habitat. The base of the productive zone of the species is on average located below 100 m and at maximum at 300 m and is regionally highly variable. The calcite flux immediately below the productive zone (export flux) is on average 8 mg $CaCO_3$ m^{-2} d^{-1}, and we observe that this flux is attenuated until at least 300 m below the base of the productive zone by a mean rate of 6.6 % per 100 m. Regionally, the summer export flux of *N. pachyderma* calcite varies by more than two orders of magnitude and the estimated mean export flux below the twilight zone is sufficient to account for about a quarter of the total pelagic carbonate flux in the region. These results indicate that estimates of the Arctic pelagic carbonate budget will have to account for large regional differences in the export flux of the major pelagic calcifiers and confirm that substantial attenuation of the export flux occurs in the twilight zone.

1 Introduction

The world's oceans play an important role in the global carbon cycle, which is at present strongly influenced by anthropogenic carbon emissions (Friedlingstein et al., 2019). With the solubility of CO_2 increasing with decreasing water temperatures, the oceanic take-up of atmospheric CO_2 is especially high in the Arctic Ocean (Steinacher et al., 2009; Miller et al., 2014). Next to the redistribution of dissolved CO_2 by ocean circulation, the surface-ocean carbon is also removed and sequestered in the deep ocean and ocean sediments by the two major carbon pumps: the biological carbon pump and the so called 'counter pump'. The biological carbon pump transports particulate organic carbon (POC) that is fixed by photosynthesis into the deep ocean where a small part of it can be buried in the sediments (Riebesell et al., 2009; Henehan et al., 2017). In contrast, the $CaCO_3$

counter pump exports biogenic carbonate produced by calcifying organisms such as pteropods, coccolithophores and planktonic foraminifera from the productive zone. Initially, CO_2 is released during calcification, but on longer time scales, a large part of the carbon fixed in biogenic carbonate is buried in the sediments and stored on geological time scales (Zeebe, 2012; Bauerfeind et al., 2014; Salter et al., 2014; Schiebel et al., 2018).

From among the pelagic calcifiers, planktonic foraminifera, calcite shell-building marine protists, are globally responsible for an estimated $CaCO_3$ sedimentation at the sea floor of 0.71 Gt yr^{-1}, accounting for more than a quarter of the global pelagic calcite flux (Schiebel, 2002). Their contribution is likely even higher in the high-latitude oceans, where the main pelagic calcite producers, the Coccolithophoridae, are less abundant (Baumann et al., 2000; Daniels et al., 2016). For example, at the Northern Svalbard margin, summertime calcite fluxes inferred from standing stocks of planktonic foraminifera at 100 m depth range from 2.3 to 7.9 mg $CaCO_3$ m^{-2} d^{-1}, which is about 4-34% of total $CaCO_3$ flux in that area (Anglada-Ortiz et al., 2021).

With ongoing global warming, the Arctic habitat is changing, becoming more hospitable for subpolar species (Wassmann et al., 2015). Pelagic calcifiers, including foraminifera, react sensitively to the ongoing transformation of their pelagic habitat (e.g. Field et al., 2006; Jonkers et al., 2019; Schiebel et al., 2018), and show increasing standing stocks in the North Atlantic (Beaugrand et al., 2013). Therefore, it is likely that continued warming and associated ecological transformation of the Arctic Ocean and its adjacent seas will also lead to changes in the carbonate counter pump and the biological carbon pump. This could have consequences for the capacity of the Arctic to take up atmospheric carbon dioxide, as well for the seawater chemistry including the nature of the sediments and thus the habitat for benthic life in this region.

In many parts of the ocean, a considerable portion of the biogenic carbonate is dissolved in the upper layer of the ocean because of processes like digestion by predators or dissolution by metabolic CO_2 released during microbial degradation of biomass surrounding the biomineral (Sulpis et al., 2021). Therefore, estimates of carbonate production and export require observations from the water column, immediately below the zone where the production occurs. Moored sediment traps provide direct observations on the seasonal cycle of biogenic carbonate flux. However, they intercept export fluxes towards the ocean floor and are typically anchored deeper than the productive zone (Wolfteich, 1994; Jensen, 1998; Jonkers et al., 2010), hence record a potentially attenuated export flux. Also, sediment trap records are too scarce in the Arctic (Soltwedel et al., 2005) to resolve the large spatial variability in planktonic foraminifera abundances and thus calcite fluxes (Volkmann, 2000b; Greco et al., 2019). Next to observations from sediment traps, planktonic foraminifera calcite fluxes can also be estimated from vertically resolved net tow profiles of standing stocks in the upper water column (Schiebel and Hemleben, 2000). Vertical profiles provide only a snapshot of the flux at the time of sampling. Also, due to the extensive sea ice cover, the time of sampling by research vessels in the Arctic is almost completely restricted to the summer season (Greco et al., 2019). However, vertically-resolved net tow profiles of shell number concentration in the water column allow us to characterise the zone in the upper water layer where carbonate production occurs and thus to quantify the new and export production as well as the rate of loss beneath it (Sulpis et al., 2021), provided that the profiles extend to below the productive zone.

The dominant planktonic foraminifera species in the Arctic Ocean is *Neogloboquadrina pachyderma* (Carstens et al., 1997; Volkmann, 2000b; Schiebel et al., 2017; Anglada-Ortiz et al., 2021). Like all extant planktonic foraminifera, the species builds its shell by sequential addition of increasingly larger chambers, such that the largest amount of calcification occurs during the final stages of its life. In addition, this species is known to often add at the end of its life cycle a calcite crust that covers all chambers of the last whorl (Kohfeld et al., 1996; Bauch et al., 1997) and can be so thick that it accounts for most of the mass of the shell (Stangeew, 2001). Encrusted specimens dominate sedimentary assemblages (Vilks, 1975; Kohfeld et al., 1996; Volkmann, 2000a), because most individuals add a crust and because encrusted shells are more resistant to dissolution.

These observations imply that understanding and quantifying the carbonate production and loss in the upper water layer by this dominant Arctic foraminifera requires understanding its vertical habitat. Many extant species of planktonic foraminifera, including *N. pachyderma*, have been suggested to perform ontogenetic vertical migration (Hemleben et al., 1989), with juvenile specimens inhabiting surface waters and slowly sinking as they age until the depth at which the last chambers or crusts are formed. Such ontogenetic migration may cause the depth where most calcification takes place to be below the abundance maximum of planktonic foraminifera. It is therefore imperative to also consider the vertical pattern of calcification. Cytoplasm-bearing specimens of *N. pachyderma* occur from the surface down to about 300 m water depth, with typically an abundance maximum around 100 m (Volkmann, 2000b; Stangeew, 2001; Greco et al., 2019). The variability of the preferred depth habitat depends on the local environmental conditions like presence of sea ice and productivity (Greco at al., 2019)

Previous work is inconclusive as to whether *N. pachyderma* performs ontogenetic vertical migration. Some studies provide evidence for an extensive ontogenetic vertical migration with the majority of calcite addition occurring towards the deep end of the habitat (Arikawa, 1983; Stangeew, 2001; Manno and Pavlov, 2014), while other studies are inconclusive (Pados et al., 2015) or indicate that calcification up to the terminal stage may occur at any depth within the habitat (Kohfeld et al., 1996; Simstich, 1999; Volkmann and Mensch, 2001). Here we make use of a large collection of vertically resolved abundance profiles of *N. pachyderma* in the Arctic and Subarctic, combining published data with new observations, to i) resolve the calcification behaviour of the species, ii) estimate its summertime calcite export flux, and iii) its attenuation below the production zone. To distinguish the production and export zones and to determine the average depth of calcification of *N. pachyderma*, we analyse vertical profiles of the abundance of cytoplasm-bearing and empty shells, shell size spectra and mean shell weights. The results allow us to constrain the spatial variability in the calcite production of *N. pachyderma* in the Arctic Ocean during summer periods, and quantify the shell dissolution within the upper water column.

2 Material and methods

2.1 Planktonic foraminifera samples

This study is based on a combination of existing and new data from vertically resolved profiles of plankton net samples from the Arctic Ocean and adjacent seas (Table 1; Fig. 1). We used all data from the studies by Kohfeld et al. (1996), Bauch et al. (1997), Kohfeld (1998), Volkmann (2000b), Stangeew (2001), Schiebel (2002), Simstich et al. (2003), Pados and Spielhagen (2014) and Greco et al. (2019), containing information on at least one of the three parameters abundance, shell size or weight/size-ratio of the planktonic foraminifera *N. pachyderma*, resulting in a data set of 112 depth profiles. As data on shell size and weight, which are important for estimates of calcite mass flux, are scarce in existing publications, we have extended the dataset by 36 new vertical profiles taken during expeditions in the Baffin Bay (MSM44, July 2015 and MSM66, July 2017) and in the Fram Strait (PS93.1, July 2015) (Table 3, Fig. 1). All of the new profiles consist of samples from five depth intervals (Table 2), sampled with a multiple closing plankton net (Hydro-Bios, Kiel) with an opening of 0.25 m² and a mesh size of 100 µm during the MSM44 and MSM66 cruises and 55 µm during PS93.1. Shell number concentrations of various planktonic foraminifera species from five depth profiles from PS93.1 are published in Greco et al. (2021b). Here we recounted the number of shells of *N. pachyderma* in those profiles, generated new counts from three further profiles in the same expedition (PS93/011-3, PS93/016-3, PS93/017-3), and added measurements on shell size and weight on shells from all eight profiles.

Samples from the Baffin Bay were either processed on board or stored at -80°C until processed onshore. All foraminifera were manually removed from each sample and counted. The counts were made separately for cytoplasm-bearing shells and empty shells, differentiated during the processing of the wet samples. As recently dead foraminifera can still contain cytoplasm, this leads to a bias in the numbers in favour of shells interpreted as being alive upon sampling. Shell size (maximum diameter) was measured with the software ImageJ on pictures taken through a SteREO Discovery.V8 microscope.

Samples from the Fram Strait were stained using a Rose Bengal/ethanol (96%) mixture to enable the differentiation of empty and cytoplasm-bearing shells. The samples were stored at 4°C until processing. They were then washed over a 250 µm and 63 µm sieve. The residues were dried on filter paper and the foraminifera were separated from the dried residues. In accordance with data from earlier studies, fully white shells were classified as empty (e.g. Fig. 2e), all other (pink) shells as cytoplasm-bearing (e.g. Fig. 2f), assumed to represent specimens that were alive during retrieval. As rose Bengal might be staining recently dead specimens because of remaining cytoplasm in the shells (Schönfeld et al., 2013), there is a possible bias towards too high numbers of cytoplasm-bearing shells. Maximum shell diameter, perimeter and area of the two-dimensional cross-section of each individual in the umbilical view were measured with a KEYENCE VHX-6000 digital microscope. As heavily calcified shells of *N. pachyderma* tend to be more round than non-encrusted specimens, the ratio of perimeter and area can indicate the foraminifera shell shape (Fig. 2 e-g): the more calcified the shell, the lower the ratio. The total weight of all shells was determined for each sample separately for shells that were considered empty and those that were considered cytoplasm-bearing, using a Sartorius SE2 ultra-micro balance (nominal resolution of 0.1 µg). The ratio between the total weight and the

mean maximum diameter (size) is here used as an indicator of the mean calcification intensity. Upon sampling, no direct differentiation between shells with or without a crust was done. ~~We regard shells as having built a crust based on their larger weight, different shell texture and more rounded shape, as exemplarily shown in Fig. 2g.~~

125 **Table 1: Overview on the used samples of vertical plankton net data of *N. pachyderma*. At M21/4 and M21/5, the profile numbers in brackets indicate the number of individually labelled and taken profiles, which were combined into fewer profiles due to sampling at the same position at different depth intervals, as indicated by the number before the brackets.**

Campaign	Region	Start Date	Number of profiles	Mesh / minimum sieving size [µm]	Reference	Data source
EN199	Greenland Sea	26.07.1989	1	150	Kohfeld (1998)	Original publication
M21/4	Norwegian Sea, Greenland Sea, Fram Strait	28.06.1992	14 (29)	100	Schiebel (2002)	from doi.org/10.1594/PANGAEA.75647 to PANGAEA.75676
M21/5	Norwegian Sea, Greenland Sea	05.07.1992	6 (14)	100	Schiebel (2002)	from doi.org/10.1594/PANGAEA.75719 to PANGAEA.75732
M39/4	Labrador Sea	12.07.1997	8	63	Stangeew (2001)	https://doi.pangaea.de/10.1594/PANGAEA.
MSM09/2	Baffin Bay	05.09.2008	8	100	Greco et al. (2019)	https://doi.pangaea.de/10.1594/PANGAEA.
MSM44	Baffin Bay	02.07.2015	13	100	This study	
MSM66	Baffin Bay	24.07.2017	15	100	This study	
NEWP-92	Fram Strait	27.07.1992	2	150	Kohfeld et al. (1996)	https://doi.pangaea.de/10.1594/PANGAEA.
NEWP-93	Fram Strait	27.07.1993	2	150	Kohfeld et al. (1996)	https://doi.pangaea.de/10.1594/PANGAEA.
ARK-IV/3 (PS11)	Nansen Basin	08.07.1987	10	160	Bauch et al. (1997)	Original publication: https://doi.org/10.1016/S0012-821X(96)00:
ARK-X/1 (PS31)	Norwegian Sea	10.07.1994	2	125	Simstich et al. (2003)	https://doi.pangaea.de/10.1594/PANGAEA.
ARK-X/2 (PS31)	Norwegian Sea	10.07.1994	1	125	Simstich et al. (2003)	https://doi.pangaea.de/10.1594/PANGAEA.
ARK-XI/1 (PS36)	Laptev Sea	03.08.1995	16	125	Volkmann (2001)	https://doi.pangaea.de/10.1594/PANGAEA.

ARK-XIII/2 (PS44)	Fram Strait, Barents Sea	27.06.1997	15	125	Volkmann (2001)	https://doi.pangaea.de/10.1594/PANGAEA.	
ARK-XV/1 (PS55)	Greenland Sea	11.07.1999	9	63	Stangeew (2001)	https://doi.pangaea.de/10.1594/PANGAEA.	
ARK-XV/2 (PS55)	Fram Strait	25.07.1999	8	63	Stangeew (2001)	https://doi.pangaea.de/10.1594/PANGAEA.	
ARK-XXVI/1 (PS78)	Fram Strait	25.06.2011	10	100	Pados & Spielhagen (2014)	https://doi.pangaea.de/10.1594/PANGAEA.	
PS93.1 (ARK-XXIX/2.1)	Fram Strait	02.07.2015	8	63	This study		

130 **Table 2: Overview on the sampled depth intervals from the stations of MSM44, MSM66 and PS93.1. Abundances of *N. pachyderma* of profiles marked with (*) are also published in Greco et al. (2021b), but counts presented in the studies were done independently from that publication.**

Campaign	Event	Longitude	Latitude	Date	Net depth intervals (m)
MSM44	MSM44/332-2	-57.982	63.074	02.07.2015	0-100, 100-200, 200-300, 300-500 500-700
	MSM44/338-2	-57.45	65.72	03.07.2015	60-80
	MSM44/339-1	-57.127	65.705	03.07.2015	0-100, 100-200, 200-300, 300-400, 400-500
	MSM44/339-2	-57.127	65.705	03.07.2015	60-80
	MSM44/340-2	-57.442	65.715	03.07.2015	60-80
	MSM44/341-2	-56.774	65.707	04.07.2015	60-80
	MSM44/348-3	-60.286	72.736	07.07.2015	60-80
	MSM44/349-1	-60.12	72.779	07.07.2015	0-100, 100-200, 200-300, 300-400, 400-500
	MSM44/349-2	-60.12	72.779	07.07.2015	60-80
	MSM44/350-2	-59.768	72.87	07.07.2015	60-80
	MSM44/351-2	-59.253	73	07.07.2015	60-80
	MSM44/355-2	-67.218	74.575	10.07.2015	0-100, 100-200, 200-300, 300-400, 400-500
	MSM44/355-3	-67.218	74.575	10.07.2015	60-80
MSM66	MSM66/4-2	-59.477	68.903	24.07.2017	0-20, 20-50, 50-100, 100-150, 150-200
	MSM66/08-2	-62.887	72.968	26.07.2017	0-20, 20-50, 50-100, 100-150, 150-200
	MSM66/13-2	-71.091	76.294	30.07.2017	0-20, 20-50, 50-100, 100-150, 150-200

	MSM66/23-2	-71.827	76.386	03.08.2017	0-20, 20-50, 50-100, 100-150, 150-200
	MSM66/27-2	-79.308	74.166	05.08.2017	0-20, 20-50, 50-100, 100-150, 150-200
	MSM66/29-2	-66.91	73.544	06.08.2017	0-20, 20-50, 50-100, 100-150, 150-200
	MSM66/33-2	-72.477	73.826	07.08.2017	0-20, 20-50, 50-100, 100-150, 150-200
	MSM66/60-3	-63.032	70	19.08.2017	0-60, 60-90, 90-120, 120-150, 150-180
	MSM66/61-2	-67	72	19.08.2017	0-20, 20-50, 50-100, 100-150, 150-200
	MSM66/61-3	-67	72	19.08.2017	0-60, 60-90, 90-120, 120-150, 150-180
	MSM66/61-4	-67	72	19.08.2017	0-60, 60-90, 90-120, 120-150, 150-180
	MSM66/62-2	-62.892	70	20.08.2017	0-20, 20-50, 50-100, 100-150, 150-200
	MSM66/63-3	-62.892	70	20.08.2017	0-30, 30-60, 60-90, 90-120, 120-150
	MSM66/63-4	-62.892	70	20.08.2017	0-20, 20-50, 50-100, 100-150, 150-200
	MSM66/65-2	-61.081	69	20.08.2017	0-20, 20-50, 50-100, 100-150, 150-200
	MSM66/65-3	-61.081	69	20.08.2017	0-30, 30-60, 60-90, 90-120, 120-150
PS93.1	PS93/011-3	-6.963	80.382	02.07.2015	0-20, 20-80, 80-140, 140-200, 200-230
	PS93/016-3	-7.341	81.217	03.07.2015	0-50, 50-100, 100-220, 220-390. 390-600
	PS93/017-3	-6.587	81.595	04.07.2015	0-20, 20-90
	PS93/020-3 (*)	-8.901	82.096	05.07.2015	0-15, 2.6-80, 80-220, 220-320, 320-600
	PS93/024-2 (*)	-6.365	80.913	07.07.2015	0-15, 15-55, 55-175, 175-350, 350-550
	PS93/030-3 (*)	-4.844	79.554	09.07.2015	0-35, 35-160, 160-250, 250-350, 350-500
	PS930/39-3 (*)	-9.612	78.748	12.07.2015	0-50, 50-150, 150-180, 180-260, 260-350
	PS93/046-2 (*)	-6.812	76.085	15.07.2015	0-75, 75-150, 150-350, 350-430, 430-500

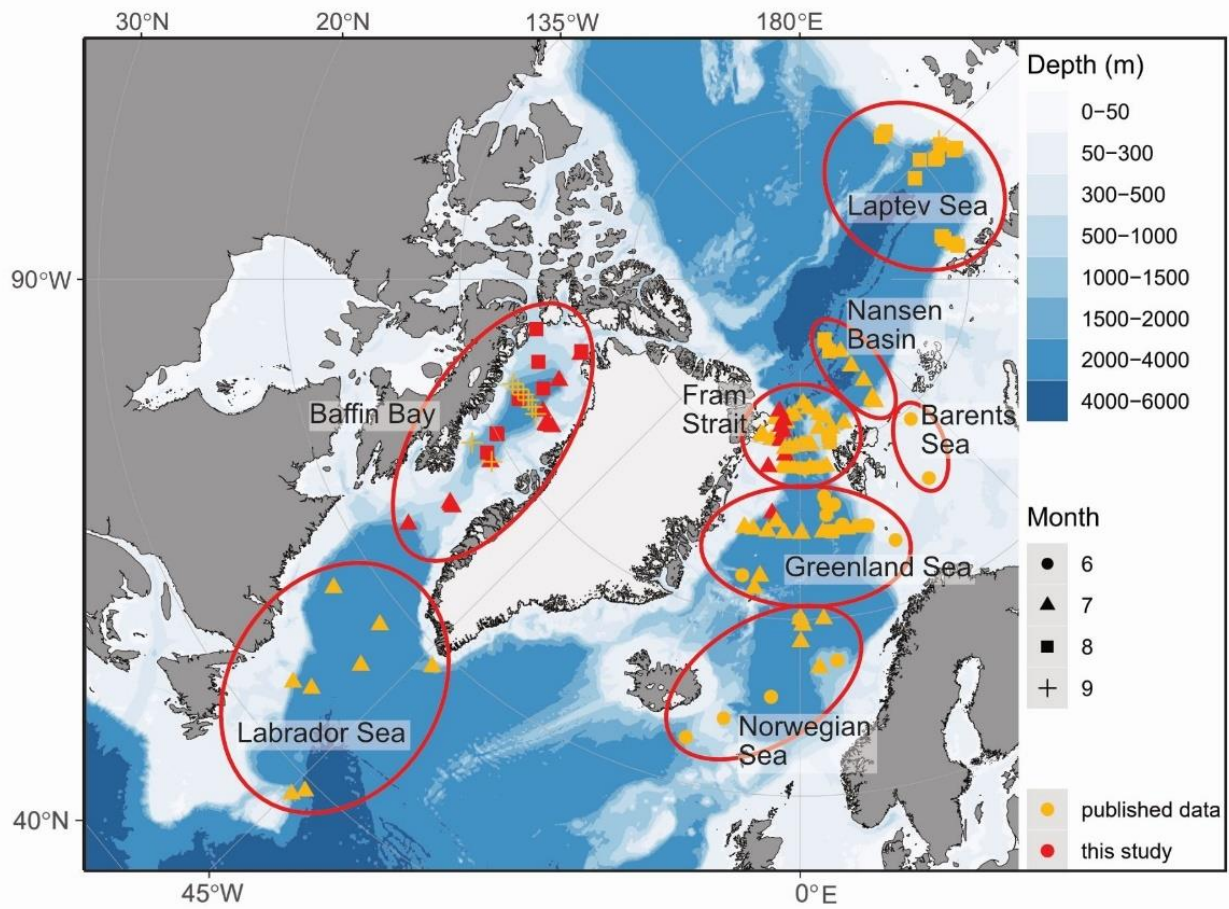



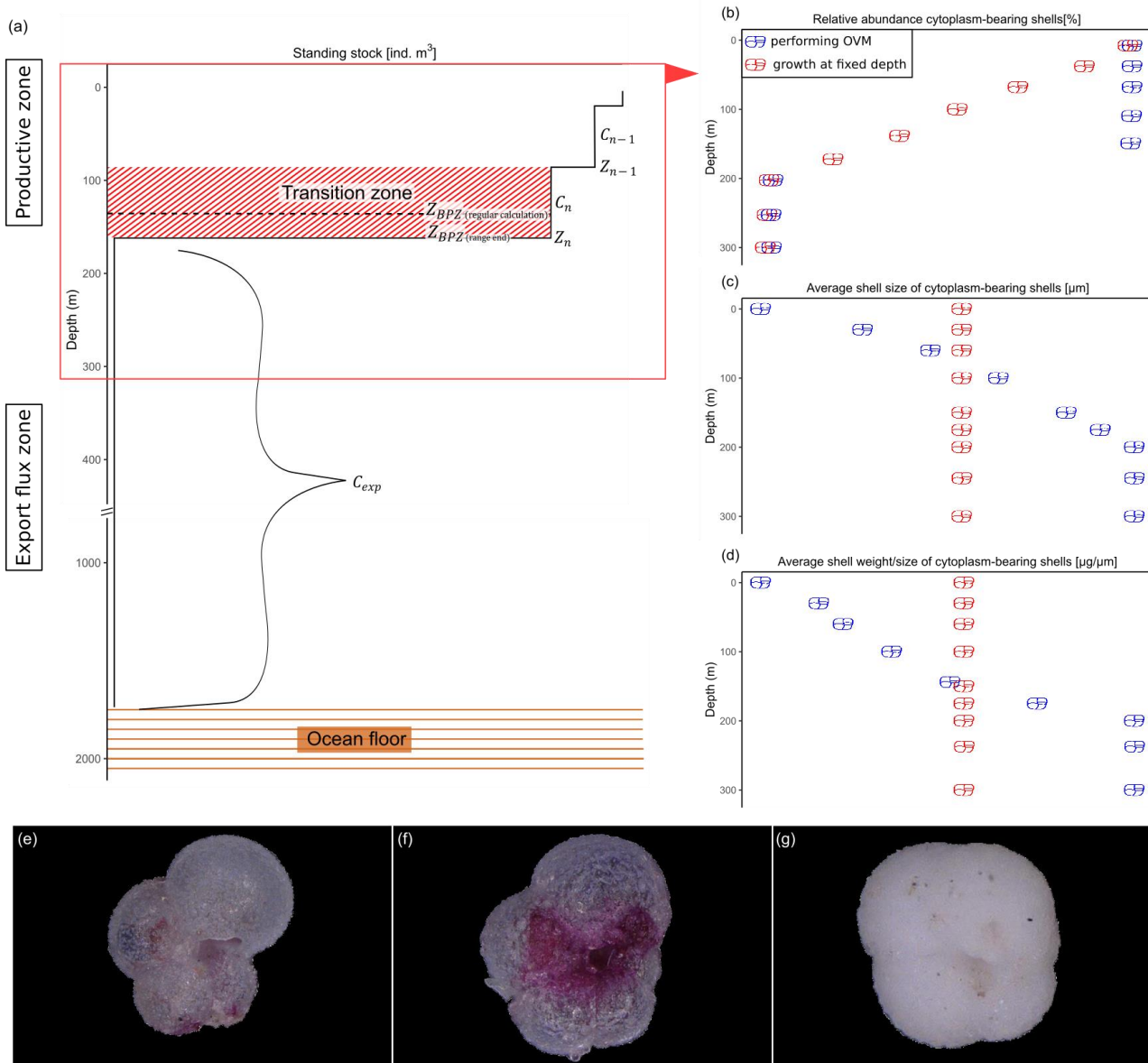
Figure 1: Overview on the research area with different regions (circled in red) sampled during different research cruises. Published data (orange) and new data sets (red) used in this study as well as the sampling periods (symbols) are marked. Land and glacier polygons from Natural Earth Data (CC0), bathymetry from Amante and Eakins (2009), using ggOceanMaps in R (Vihtakari, 2021).

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Figure 2: Schematic overview on the studied shell parameter. Shown values are constructed numbers to represent the concept of the study and not measured values. (a) change of standing stock of planktonic foraminifera with increasing depth. The parameters used to calculate the base of the export zone (Z_{BPZ}) after Lončarić et al. (2006) are shown: The transition zone represents the area in which the foraminifera shell abundance (C_n) rapidly changes, with rather stable abundances in the area below (C_{exp}). Z_n and Z_{n-1} represent the start and end depth of the transition zone, in which the calculated BPZ is located. For details on the calculation, see Sect. 2.2. (b, c, d) show the change of average (b) relative abundance of cytoplasm-bearing shells, (c) average shell size and (d) average calcification intensity (shell weight/size) of cytoplasm-bearing shells with increasing water depth within the productive zone. Blue symbols represent the ideal situation if *N. pachyderma* performs ontogenetic vertical migration (OVM) throughout its lifecycle, while red shell symbols indicate the expected trend when individual specimens grow their shell at a fixed depth. (e, f, g) show different types of encrustation of *N. pachyderma*, with (e) representing a non-encrusted shell, (f) the beginning of encrustation and (g) thick encrustation with a clearly different and more rounded shape.

2.2 Productive Zone

150   To determine the depth range where shell calcification occurred and below which the export began, the base of the productive zone (BPZ) of *N. pachyderma* was defined for each profile by considering the changes in shell abundance with depth. Following the concept of Peeters and Brummer (2002), the BPZ is the depth where the shell abundance begins to substantially decline. It was calculated after Lončarić et al. (2006):

$$Z_{BPZ} = \frac{C_n - C_{exp}}{C_{n-1}}(Z_n - Z_{n-1}) + Z_{n-1} \qquad (1)$$

155   where C_n is the concentration of shell numbers within the transition zone (i.e. the last depth interval before the rapid decline in shell abundance) which was defined visually for every profile as exemplarily shown in Fig. 2a,

C_{exp} is the average shell abundance, weighted by the thickness of the sampled depth interval, in all depths below C_n,

C_{n-1} is the foraminifera abundance in the depth interval above C_n. Z_n represents the top of sampling depth of the transition zone, and Z_{n-1} its bottom.

160   The equation applies to cases where the shell number concentration decreases with depth. Where this is not the case (such as where there is a distinct subsurface maximum), the equation cannot be used as the estimated BPZ would appear to lie below the depth interval of the transition zone. This was the case in 37 out of 126 profiles. In addition, in three profiles, the transition zone corresponded to the uppermost sampling layer, and the equation could not be applied. For those 40 profiles, the BPZ was defined as the bottom depth of the transition zone (Fig. 2a, $Z_{BPZ\ (range\ end)}$). This can result in a bias towards the estimated BPZ

165   being located below the actual position. This bias is restricted by the overall sampling interval (median: 50 m) and has no effect on our flux estimates which are based on average shell abundances below the BPZ. In ten profiles, calculation of the BPZ was not possible as no clear transition zone was present within the sample range, including two profiles in which the abundance was zero at the total station. The maximum sampling depth of those profiles was between 180 and 300 m, implying that the transition zone either occurred in the bottom interval or was not yet reached. Because of this ambiguity, these profiles

170   were not used for the BPZ analysis. For profiles where abundance data were available for only one or two depth intervals at the surface (9 Profiles), estimation of the BPZ was not possible either. In total, the BPZ was determined in 126 profiles and the different methods to define BPZ were separated in the interpretation. For an overview on the number of profiles that were available for the different calculations, see table 3.

The above definition of the BPZ does not rely on the separation of living (cytoplasm-bearing) and dead (empty) shells during

175   sampling, a parameter that was not systematically recorded. The separation is ambiguous as cytoplasm decomposition takes time after death and shells already dead could still be considered as living due to the presence of residual cytoplasm (Schiebel et al., 1995). This ambiguity is larger at greater depth, where the probability of finding living specimens becomes smaller.

Nevertheless, where available, we used the proportion of cytoplasm-bearing and empty shells as another indicator of the maximum extent of the productive zone.

180   To investigate at which depth of the productive zone the calcification of *N. pachyderma* occurred and if the species performed ontogenetic vertical migration, we considered the vertical profiles of the following parameters: (i) relative abundance of empty shells, (ii) shell size and (iii) mean calcification intensity expressed as the shell weight/size-ratio. The reason for using those parameters is that if *N. pachyderma* performed ontogenetic vertical migration and premature mortality were zero, empty shells would only be present at the bottom of the productive zone, where the specimens would reach their maturity, while the
185   abundance of cytoplasm-bearing shells would be 100 % at all depths above (Fig. 2b). At the same time, shell size and calcification intensity would increase constantly with increasing depth, reaching maximum values only at the base of the productive zone. In contrast, if individual specimens did not migrate during their life cycle, the fraction of the population dying would be equal across the productive zone. Assuming that empty shells only sink, this would lead to a linear decrease in relative abundance of cytoplasm-bearing shells. Because foraminifera of any life stage would be present in equal proportions
190   at all depths, the average shell size and weight of cytoplasm-bearing specimens should stay constant with increasing depth (Fig. 2b-d).

Table 3: Overview on the numbers of depth profiles used in the study, with varying numbers depending on the studied parameter.

Total profiles		148
	published data	112
	new profiles added by this study	36
Profiles to determine BPZ		126
	calculated after Lončarić et al. (2006)	86
	determined by range end	40
Profiles with size measurements		23
Profiles with calcification intensity measurements		13
Profiles with calcification intensity trend		9
	cytoplasm-bearing shells	6
	empty shells	6
	non-encrusted shells	3
	(heavily) encrusted shells	3
Profiles to calculate mass flux		147

2.3 Export flux zone

When the bottom of the productive zone is known (or estimated), the abundance of shells below that depth can be used to estimate the export flux by taking the sinking velocity into account (Schiebel and Hemleben, 2000). Assuming that the organic matter content of foraminifera is negligible, the calcite flux can subsequently be calculated using (average) shell weight

Calcite mass flux = average shell weight * shell number concentration * sinking velocity (2)

where shell weight is the measured average weight of shells below the productive zone, as these are representative of the export flux. Whenever possible, the measured average shell weight was used, but for samples where no weight data is available, we used regional mean values. In regions where some weight data were available (Fram Strait, Labrador Sea, Greenland Sea, Norwegian Sea), average weights were calculated from samples of those regions alone. In all other regions, the overall mean weights from our data were used. This method is likely to underestimate present variability. To evaluate possible effects on mass flux from distinct shell types, fluxes based on average weights of either only encrusted and empty or non-encrusted and cytoplasm-bearing shells from below the productive zone were calculated as well. Shell abundance was calculated as the number of shells, divided by the sampled depth range and multiplied by the area of the net opening (as an estimate for the volume of filtered water). Sinking velocity was calculated after Takahashi and Bé (1984):

Foraminifera sinking velocity (m d^{-1}) = $10^{2.06}$ * shell weight $^{0.64}$ (3)

using the same (average) weights as described above.

The residence time of *N. pachyderma* in the productive zone was then estimated based on the standing stock within the productive zone (ind. m^{-2}) divided by the shell flux (ind. m^{-2} d^{-1}).

2.4 Statistical analysis

Statistical analyses were performed using R v. 3.6.1 (R Core Team, 2018). To compare measured parameters between cytoplasm-bearing and empty shells, a Welch's t-test was performed. The analysis of trends within the productive zone was done within the beforehand individually calculated range of the productive zone of the stations. Linear regression models were used to detect the effects of depth and sampling location on the different parameters. As the data of shell size and calcification intensity is not normally distributed, it was log-transformed before these analyses. Since the depth of the BPZ varies among the profiles, analyses were performed using tow intervals standardised to the depth of the productive zone. Some intervals extend to below the BPZ. In these cases, the tow interval represents >100 % of the depth of the productive zone.

3 Results

3.1 Shell abundances and the productive zone

The average shell abundance of *N. pachyderma* in our dataset is 25 ind. m^{-3} (Table 4). Shell abundances show either a maximum within the upper 50 m, or in the depth zone below, reaching down to 150 m (exemplarily shown in Fig. A1). Those distinct patterns are distributed rather equally among all profiles and regions. Below the depth of maximum shell abundance, there is a rapid decrease in all profiles, until the abundances stabilise~~, which occurs no deeper than at around 300 m~~. Empty shells of *N. pachyderma* are present across the entire sampled depth range (Fig. 3a). In the majority of the profiles, the BPZ is located between 100 m and 150 m (Fig. 3b). The median BPZ based on the calculation after Lončarić et al. (2006) is situated at 124 m. At stations where the BPZ could only be defined as the end of the depth range of the transition zone, the median is 136 m. Irrespective of how calculated, the BPZ varies among different stations and regions, with the lowest median ~~value~~ of 100 m in the Baffin Bay, and the highest median ~~value~~ of 160 m in the Barents Sea (Table 4), with variability within the regions being as large as among the regions. The minimum calculated BPZ is 15 m in a profile from the Fram Strait (PS93/020-3) and the minimum BPZ determined by the end of the net range is 20 m in a profile from the Baffin Bay (MSM09/2 466-2). The deepest BPZs reach 300 m and ~~are in line~~ with the pattern visible in the relative abundance of empty shells (Fig. 3a). Within the productive zone, the average shell number concentration of *N. pachyderma* is 42.27 ind. m^{-3}, below the productive zone, it is 6.52 ind. m^{-3} (Table 4).

Table 4: Overview on measurements on different shell parameters from the different sampling areas of the study. Next to each indicated value, the 95% confidence interval (CI) is given in italics. It is calculated assuming normal distribution, and the number of samples (n) used to calculate each parameter is given in brackets.

	Shell abundance (ind. m-3)						BPZ (m)		Shell size (µm)			Shell weight (µg)			Mean mass flux (mg CaCO3 m-2 d-1)					
	Mean	*95% CI (n)*	Mean productive zone	*95% CI (n)*	Mean export flux zone	*95% CI (n)*	Median	*95% CI (n)*	Mean	Median	*95% CI (n)*	Mean	Median	*95% CI (n)*	100m	*95% CI (n)*	Below BPZ	*95% CI (n)*	Deepest sampling position	*95% CI (n)*
Arctic (all samples)	25.4	*4.3 (825)*	42.3	*6.1 (404)*	6.5	*0.0 (360)*	112.7	*10.8 (127)*	150.0	143.4	*0.5 (40275)*	3.4	2.3	*0.3 (330)*	40.8	*10.3 (173)*	8.0	*3.0 (126)*	4.4	*2.3 (126)*
Baffin Bay	50.9	*11.9 (176)*	90.4	*24.0 (71)*	16.4	*6.6 (77)*	100.0	*28.3 (22)*	146.5	140.6	*0.5 (36070)*				85.5	*26.1 (44)*	22.7	*14.3 (21)*	13.7	*12.8 (21)*
Barents Sea	0.6	*0.3 (10)*	0.6	*0.4 (7)*	0.5	*0.3 (3)*	160.3	*51.8 (2)*							0.7	*0.6 (2)*	0.5	*0.5 (2)*	0.5	*0.5 (2)*
Fram Strait	11.3	*2.0 (265)*	19.5	*3.6 (126)*	3.2	*0.9 (120)*	100.0	*16.7 (44)*	180.0	172.1	*1.6 (4205)*	3.0	2.1	*0.4 (193)*	18.3	*5.3 (50)*	3.9	*1.6 (44)*	1.6	*0.9 (44)*
Greenland Sea	45.2	*19.7 (110)*	79.2	*36.5 (56)*	9.9	*4.3 (45)*	116.1	*27.7 (14)*				3.0	3.0	*0.5 (40)*	66.6	*45.4 (23)*	17.1	*10.6 (14)*	8.0	*4.1 (14)*
Labrador Sea	8.7	*6.3*	14.2	*11.5*	2.6	*1.5*	15	*59.7*				4.4	2.8	*0.8*	6.6	*3.4*	2.4	*2.0*	4.3	*4.8*

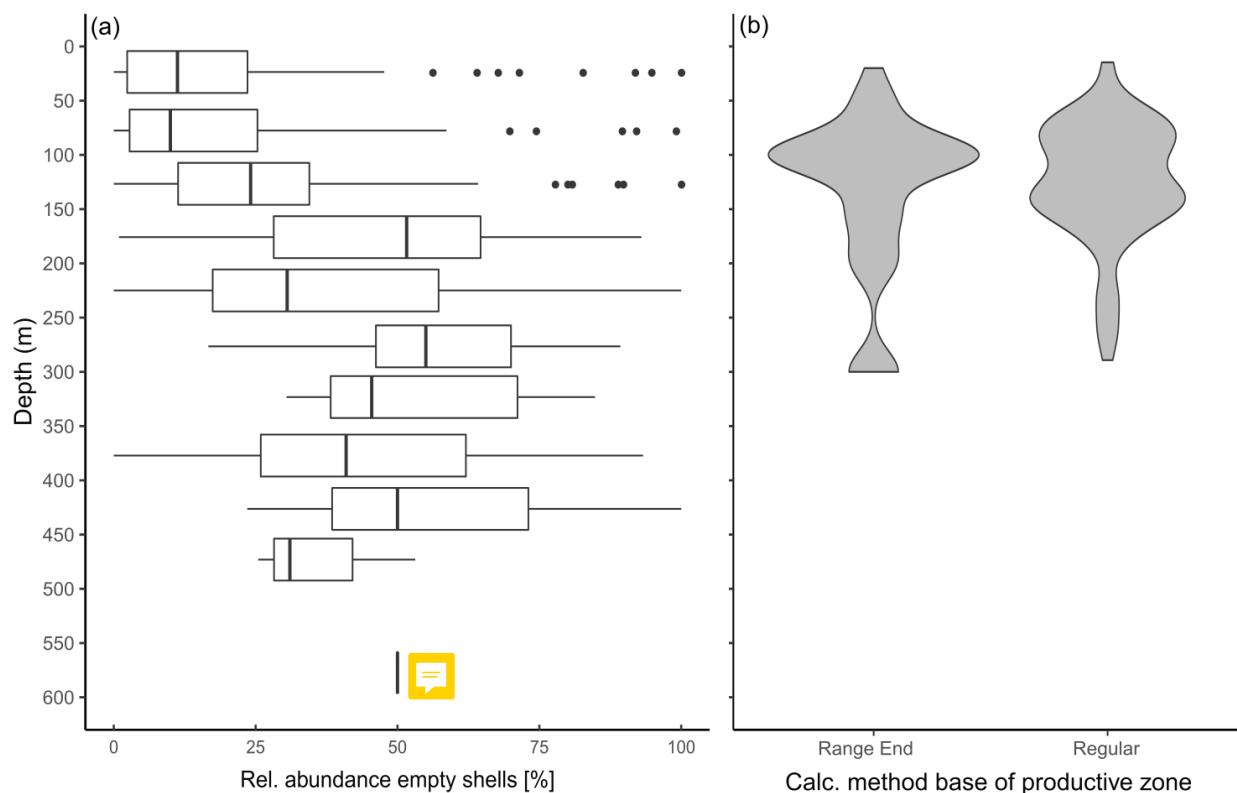
		(38)		(20)		(18)	3.0	(8)				(97)	(8)		(8)			(8)
Laptev Sea	2.0	0.5 (81)	2.8	0.8 (41)	0.7	0.2 (35)	139.7	31.9 (15)					4.3	1.7 (16)	1.0	0.4 (15)	0.5	0.4 (15)
Nansen Basin	20.0	5.9 (50)	35.8	7.1 (26)	2.9	0.9 (24)	106.4	36.8 (10)					34.8	7.9 (10)	5.1	1.2 (10)	2.3	1.6 (10)
Norwegian Sea	26.8	14.4 (95)	42.7	23.2 (57)	2.8	1.0 (38)	125.8	51.4 (12)					19.1	8.8 (20)	3.0	2.5 (12)	1.8	1.3 (12)

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Figure 3: (a) Vertical Profile of relative abundance of empty shells at all stations of the study in which empty and filled shells were distinguished. The boxes represent the interquartile range (IQR) of the relative abundance at the given depth, and the vertical bar represents the median. Outliers, shown as points, are values beyond 1.5*IQR of each site of the box, and lines represent the range within 1.5*IQR. (b) Range of the base of the productive zone (BPZ), divided by the way they were determined: "Range end" shows all samples in which the maximum depth of the net of the transition zone was defined as the BPZ, while "Regular" shows all samples in which the equation from Lončarić et al. (2006) to estimate BPZ could be applied, as described in Sect. 2.2.

3.2 Shell sizes

The average maximum diameter of *N. pachyderma* in our samples is 150 µm (Table 4). Shells from the Baffin Bay with a mean size of 146.5 µm (sampling mesh size: 100 µm) are smaller than shells from the Fram Strait (only data from PS93.1) that have a mean size of 180 µm (sieving size: 63 µm; Table 4, Fig. 4). A Welch's t-test shows that this difference is significant ($p < 0.001$). Cytoplasm-bearing shells within the estimated productive zone of each station in samples from PS93.1 are on average bigger than empty ones (mean sizes of 188.2 µm and 166.2 µm, respectively; Fig. 4a). A Welch's t-test shows that

this difference is significant in eight of 14 individual samples (p ≤ 0.006). At station PS93/024-2 in the topmost net (0-15 m), empty shells were significantly bigger (p = 0.035) than cytoplasm-bearing ones. Below the productive zone, two of 16 individual sampling positions contain empty shells that are on average significantly bigger than those filled with cytoplasm (p < 0.01). In all other samples, the differences were not statistically significant. In both regions, shells below the productive

270 zone are significantly, if only slightly, bigger than within the productive zone (Welch's t-test: p < 0.001), with averages of 150 µm and 153 µm, respectively (Fig. 4b). Statistical analysis indicates that there is no significant linear increase in average size within the productive zone (Fig. 5, Baffin Bay: p = 0.399, Fram Strait empty: p = 0.199, Fram Strait cytoplasm-bearing: p = 0.627). We find no evidence for lunar periodicity in the shell size of *N. pachyderma* in our samples.

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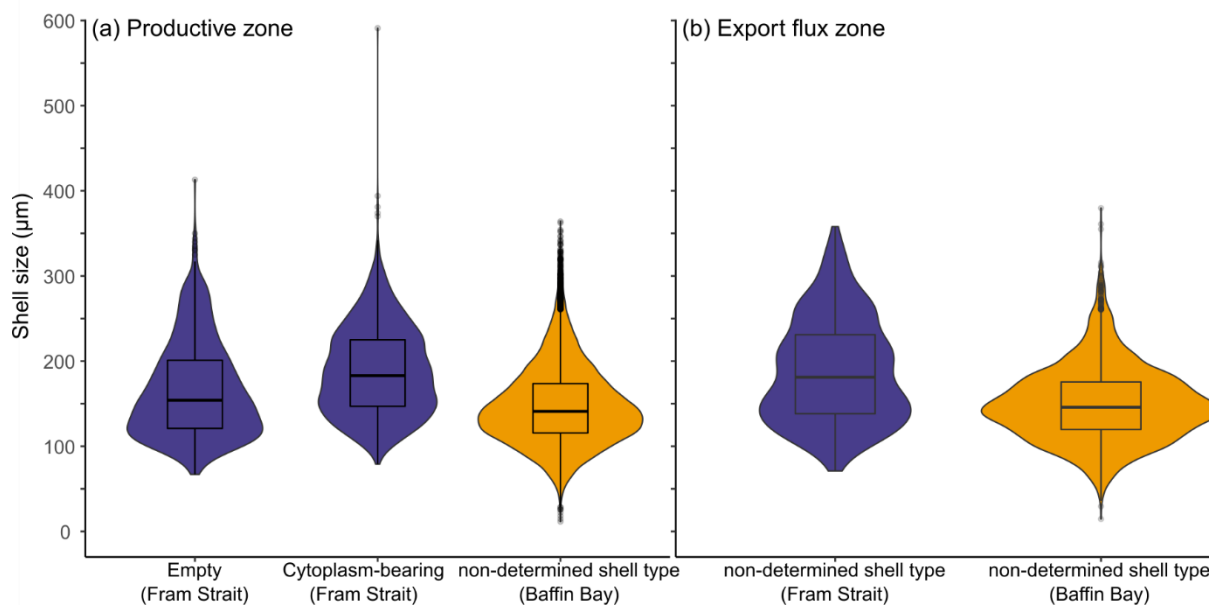


Figure 4: Overview of shell sizes of *N. pachyderma* from the Fram Strait (blue) and the Baffin Bay (orange), contrasting empty, cytoplasm-bearing and non-determined shells (a) within the productive zone, and (b) in the export flux zone. Shell types are also not
280 **distinguished in (b) in samples from the Fram Strait as we assume all shells collected below the productive zone to represent specimens that were dead during retrieval. The boxes and bars represent the interquartile range as explained in the caption of Fig. 3.**

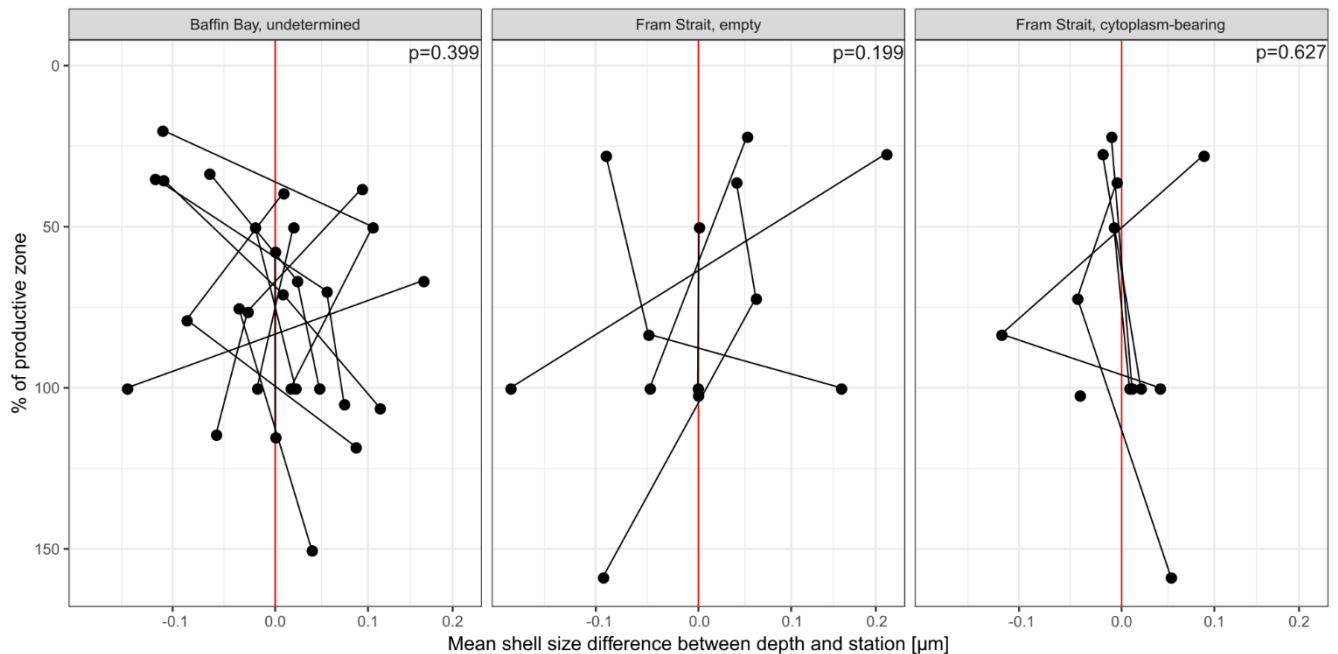



Figure 5: Mean of difference in mean shell size at the individual station and depth and the overall mean of the station, plotted against the percentage of the depth interval on the overall depth of the productive zone. 100 % equals the total depth of the productive zone, 50 % half of the depth of the productive zone. More than 100 % are reached where the sampling interval ends below the BPZ. The plot is divided into different types of shells (undetermined, empty, cytoplasm-bearing) and the two regions from which size measurements are present (Baffin Bay, Fram Strait). Consider that the samples do not represent all samples from the region shown in Fig. 1, but only those from PS93.1 (Fram Strait), MSM44 and MSM66 (Baffin Bay). The red line indicates the position at which no difference between the mean of the depth and the overall station exists. Only the depth interval within the estimated productive zone of each station is shown. p-values show the effect of increasing proportion of productive zone on shell size.

3.3 Shell calcification intensity

Across both new and literature data, the mean shell weight of *N. pachyderma* per sample ranges from 0.1 µg (potentially referring to fragments of shells from the Fram Strait, data from Kohfeld, 1998) to 20.8 µg (shells from the Labrador Sea, data from Stangeew, 2001). The overall average weight is 3.4 µg (median: 2.3 µg, Table 4) and the average calcification intensity (weight/size) 0.013 µg/µm (median 0.010 µg/µm). A clear difference in weight and calcification intensity is present between (heavily) encrusted and non-encrusted shells, with the latter being lighter and having a lower calcification intensity. Similarly, cytoplasm-bearing shells are lighter and have a lower calcification intensity than empty shells (Fig. 6). The differences become smaller below the productive zone. A Welch's t-test shows a significant difference between the calcification intensity of cytoplasm-bearing and empty shells from PS93.1, both within (p < 0.001) and below (p = 0.004) the productive zone, with empty shells being always stronger calcified.

Shell size parameters can be used to infer the presence of crust by a less lobate periphery (see Fig. 2 e-g) in samples where it
305 has not been checked visually: Lower perimeter-area-ratios indicate rounder, likely more encrusted, shells. Indeed, both within
and below the productive zone, empty shells from PS93.1 are significantly rounder than cytoplasm-bearing shells (Welch t-
test, $p < 0.001$), suggesting that empty shells are more encrusted than cytoplasm-bearing shells (Fig. A2). We observe no
statistically significant difference in the roundness of shells between cytoplasm-bearing shells within and below the productive
zone ($p = 0.9$), but empty shells from below the productive zone are significantly rounder than those within the productive
310 zone ($p < 0.001$; Fig. A2). While differences within samples from the Fram Strait could be partly due to differences in sampling
methods among the different studies and authors, large regional differences between the Fram Strait, the Greenland Sea and
the Labrador Sea are likely reflecting real variability, because many of the involved studies used the same methodology (Fig.
6).

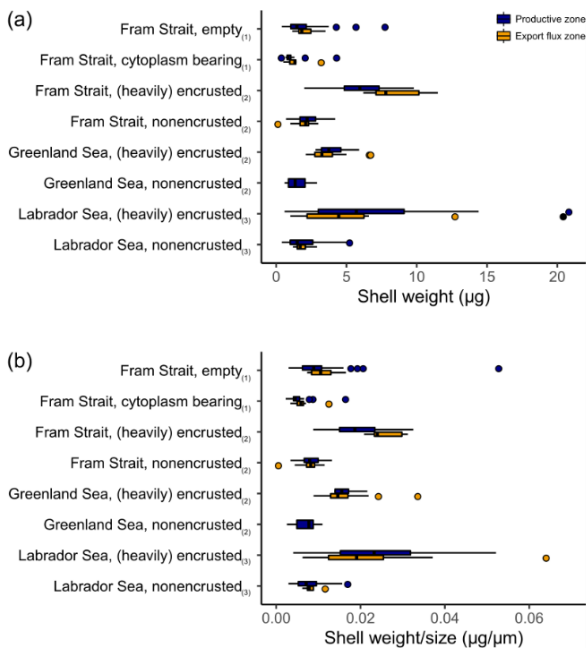
315 Ten out of 18 profiles show a clear tendency towards higher calcification intensity with depth (Fig. 7). In seven profiles, no
clear trend with depth can be detected in calcification intensity. Those profiles are all from the samples of PS93.1, four of them
of empty and three of cytoplasm-bearing shells. One profile of non-encrusted shells from the Fram Strait shows lower
calcification intensity at deeper depth. The involved sample size is too small to allow statistical analysis.

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**Figure 6: Overview on average shell weight (a) and calcification intensity (weight/size) (b) from shells with a different status. In this study (1), determination was made between cytoplasm-bearing and empty shells on shells ≥ 63 µm, while Kohfeld (1998; shells ≥150 µm; (2)) and Stangeew (2001; shells ≥ 63 µm; (3)) distinguished between (heavily) encrusted and non-encrusted shells. Besides, different sampling regions are distinguished. Blue boxes show the parameter within the productive zone of each station, orange
325 boxes the values from samples taken below the estimated productive zone of each station. The boxes and bars represent the interquartile range as explained in the caption of Fig. 3.**
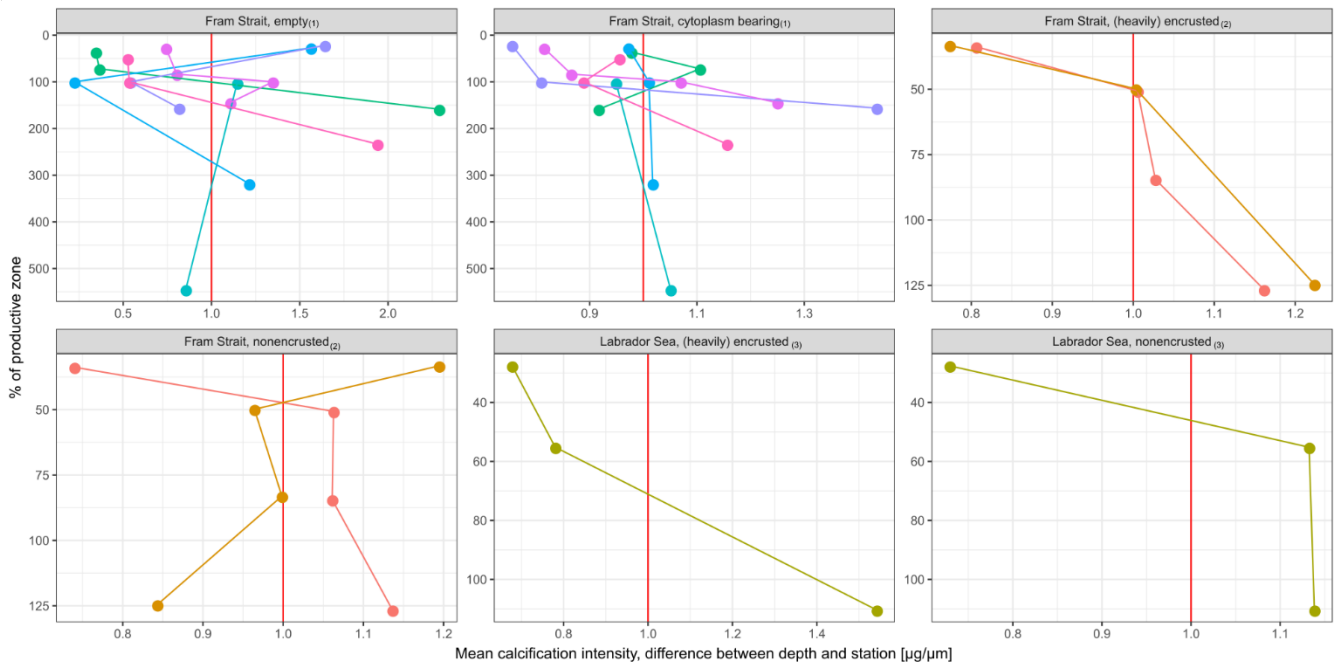
Figure 7: Mean of the difference in average calcification intensity (weight/size) at individual stations and depths and the overall weighted mean of each station within the productive zone, plotted against the percentage of the depth interval on the overall depth
330 **of the productive zone. 100 % equals the total depth of the productive zone, 50 % half of the depth of the productive zone. More than 100 % are reached where the sampling interval ends below the BPZ. Differentiation of shell types is done between cytoplasm-bearing and empty shells from Fram Strait samples of this study (1), while Kohfeld et al. (1998) (2) and Stangeew (2001) (3) distinguished between (heavily) encrusted and non-encrusted shells in samples from the Fram Strait and the Labrador Sea. The red line indicates the position at which no difference between the mean of the depth and the overall station exists, different colours are**
335 **used to make the shape of change in individual profiles visible.**

3.4 Shell mass flux

The overall mean calcite mass flux of shells of *N. pachyderma* below the BPZ in each profile based on actual weights or, where not measured, average weight of shells from below the productive zone, is 8.0 mg $CaCO_3$ m^{-2} d^{-1} (20.1 mg $CaCO_3$ m^{-2} d^{-1} based on weights of encrusted/empty shells only; 4.5 mg $CaCO_3$ m^{-2} d^{-1} based on weights of non-encrusted/filled shells only; in the

340 following, those two values will always be given in brackets without further stating this specification). Although in some profiles, the flux seems to increase further below the BPZ, the majority of the profiles shows almost no change or a decrease in mass flux (Fig. 8). When calculated for shell number concentrations at the deepest net of each profile, the average calcite mass flux is further reduced by a half to 4.4 mg $CaCO_3$ m^{-2} d^{-1} (10.7 mg $CaCO_3$ m^{-2} d^{-1}; 2.4 mg $CaCO_3$ m^{-2} d^{-1}). The average loss rate in fluxes of $CaCO_3$ from the net below the base of the productive zone and the deepest sampling position of each

345 profile is 6.6 %/100 m (8.9 %/100m; 9.5%/100 m), the median loss is 9.1 %/100 m (19.4 %/100m; 19.4 %/100 m). The highest variations and most extreme values of changes with depth are present in the Baffin Bay, the Fram Strait and the Labrador Sea (Fig. A3). Scaling the calcite mass loss for every pair of depth intervals below the BPZ (Fig. 9b) reveals that high values (and high variability of values) are limited to the 100 m below the BPZ, with both mean values and variability decreasing with

depth. Weight measurements from the profiles of PS93.1 indicate that this loss is both driven by a decrease in shell mass and shell number concentration (Fig. 9a).

Irrespective of how (at which depth) the flux was calculated, the estimated mass fluxes varied among the 148 profiles by more than three orders of magnitude (Fig. 10). This variability has some regional components: the highest flux below the productive zone (156.9 mg $CaCO_3$ m^{-2} d^{-1}; 398.6 mg $CaCO_3$ m^{-2} d^{-1}; 83.4 mg $CaCO_3$ m^{-2} d^{-1}) was determined for a station in the central Baffin Bay (Fig. 11a). In the Greenland Sea, some stations also show high values (maximum of 66.64 mg $CaCO_3$ m^{-2} d^{-1} based on individual measurements). Those two regions are also the regions with the highest average fluxes (both about 20 mg $CaCO_3$ m^{-2} d^{-1} at the base of the productive zone based on individual measurements). In comparison, average fluxes are low in the Barents Sea, Fram Strait, Labrador Sea, Laptev Sea and Norwegian Sea (< 5 mg $CaCO_3$ m^{-2} d^{-1}, Table 4). The spatial distribution of relative abundances of *N. pachyderma* in sedimentary assemblages (Siccha and Kucera, 2017) reveals that the estimated calcite fluxes from *N. pachyderma* alone from the Laptev Sea, the Nansen Basin and the Baffin Bay likely represent total foraminifera calcite fluxes, whereas in the other regions other species contribute more to the foraminifera calcite flux (Fig. 11b).
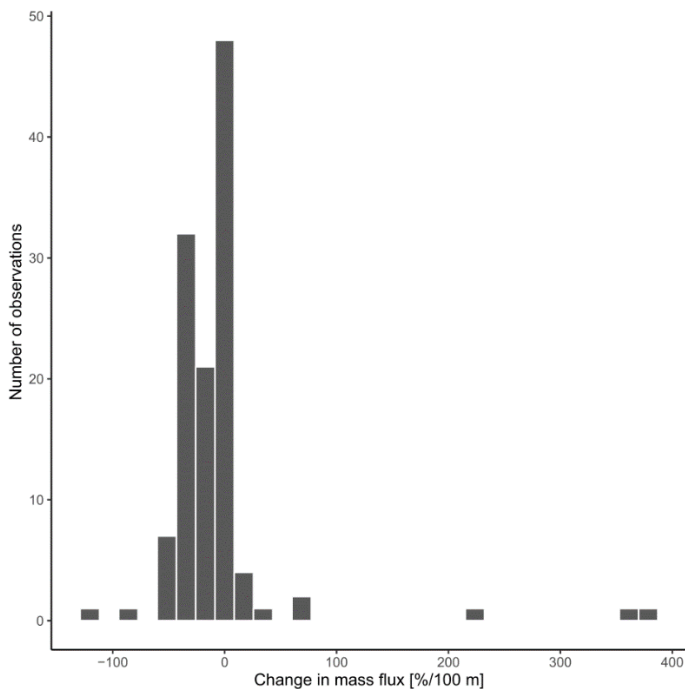
Figure 8: Loss in shell flux between the net directly below the calculated base of the productive zone and the deepest net of sampling of each station.
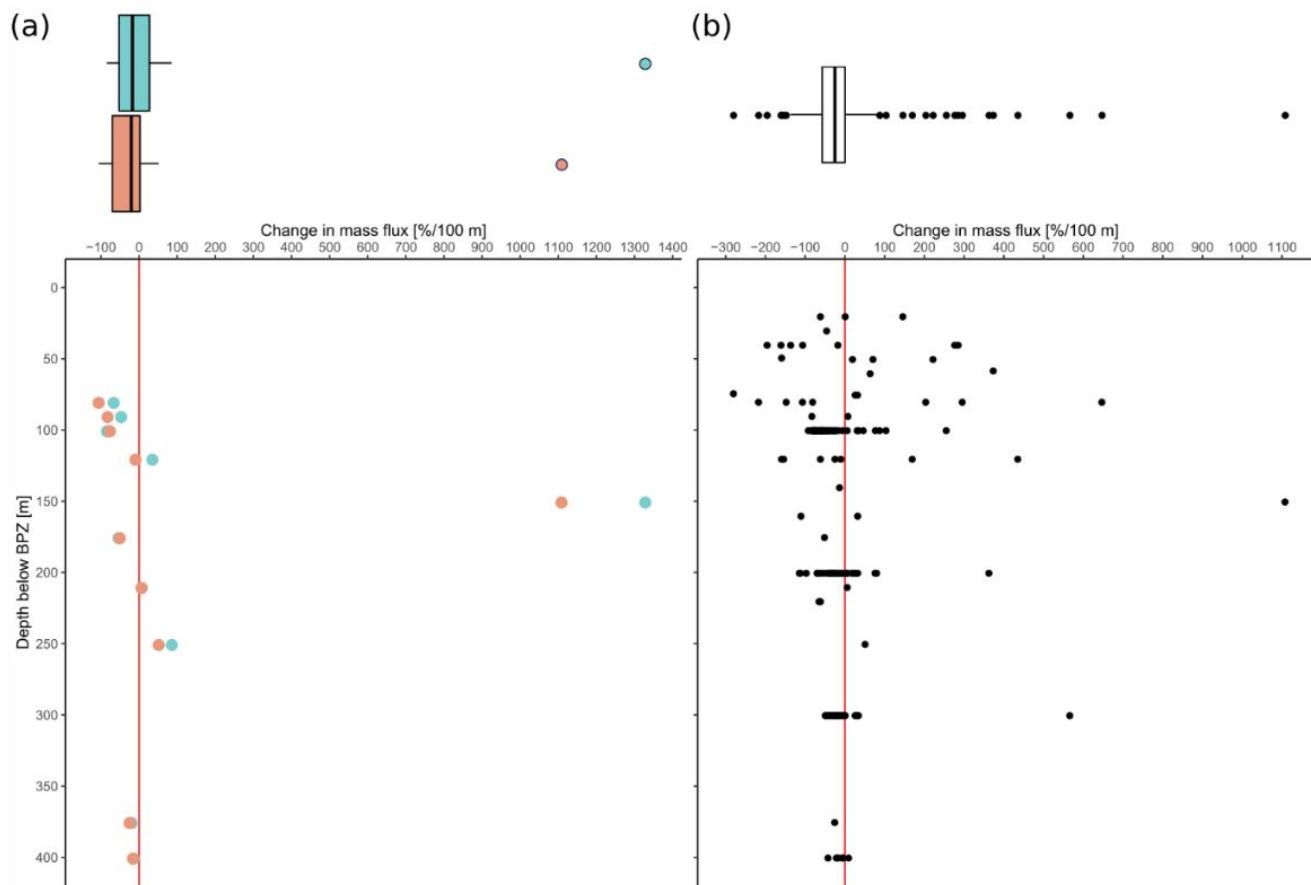
Figure 9: Flux loss with depth per 100 m in %, calculated between different sampling intervals located below the interval including the base of productive zone, plotted against the distance between the maximum sampling depth of the individual interval and the end of the net including the base of the productive zone. (a) is a comparison of loss in shell number concentration (blue) and shell mass (orange) in PS93.1 samples from the Fram Strait, (b) shows the loss in mass flux at all samples, estimated based on average shell weight and shell number concentration. The boxes and bars on top of the plots represent the interquartile range as explained in the caption of Fig. 3 and are plotted against the same x-axis as the plot below.

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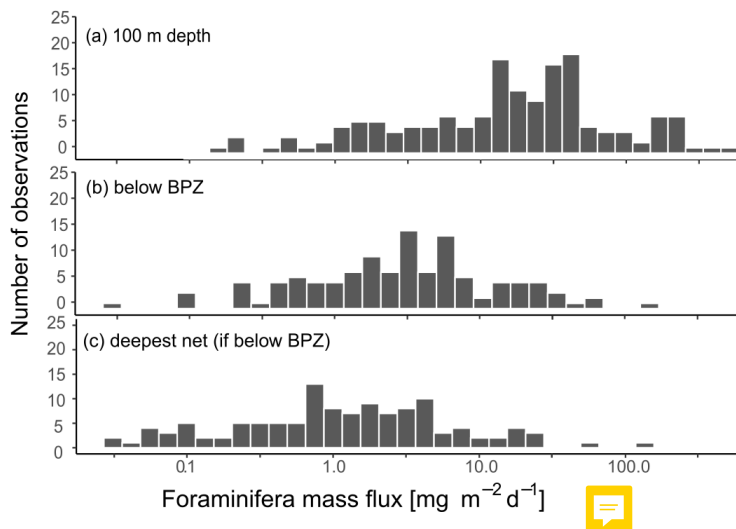
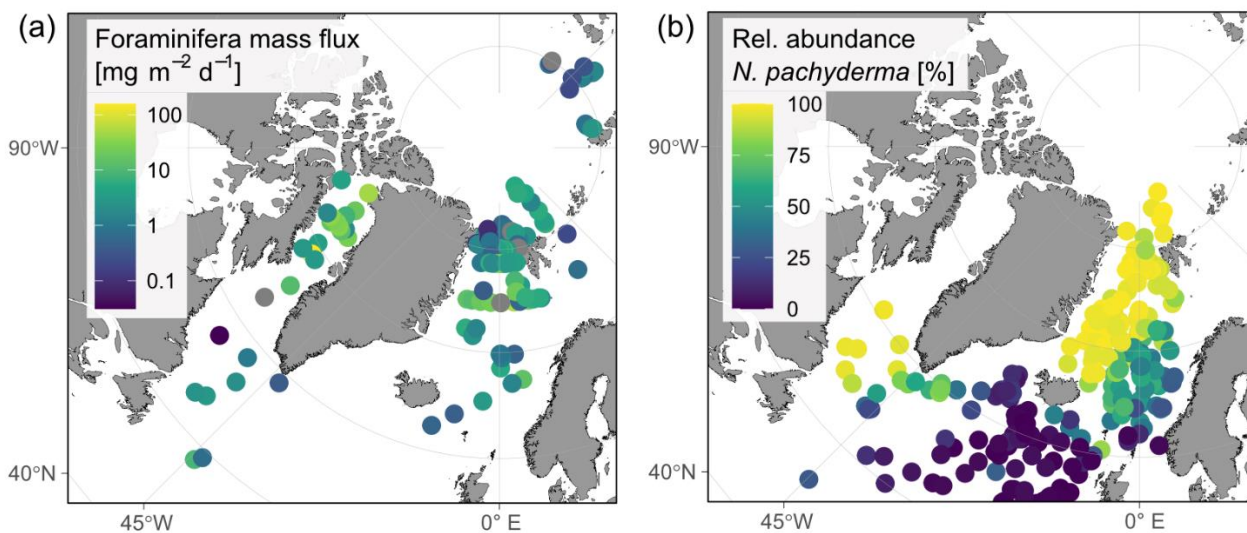

375

Figure 10: **Flux** of planktonic foraminifera *N. pachyderma*, calculated based on shell weights of individual samples, and, where no weight measurements are present, based on average weights from the region or all samples included in this study. Consider the logarithmic scale of the x-axis. (a) shows the fluxes at around 100 m depth (maximum sampling depths of nets: 75-100 m), (b) the

380 flux in the net below the calculated base of productive zone (BPZ) of the individual stations and (c) at the deepest net of each station including all stations where it is located below the BPZ. The exact width of sampling intervals differs between individual sampling locations. Details on this are shown in Tbl. 2 for profiles added in the study, and in the linked references listed in Table 1.

385 Figure 11: Regional overview on (a) foraminifera **mass** flux of planktonic foraminifera *N. pachyderma* during summer (sampling period from June to September, varies among stations) below the estimated productive zone. Fluxes were calculated based on shell abundances determined in plankton net samples. Shell weights are either from direct measurements or based on average weights from the region of sampling. Consider that values are plotted on a logarithmic scale to visualise the huge regional variability. **(b) shows the relative abundance of the species *N. pachyderma* found in sediment cores (data from ForCenS data set, Siccha and Kucera,**

390 **2017).**

3.5 Residence time

The calculated residence time of *N. pachyderma* based on standing stock within and shell fluxes below the productive zone ranges from < 1 to 79 days, including three extreme values of 182 (MSM09/2 455-7, Baffin Bay), 373 (M21/4 MSN697 and MSN698, Norwegian Sea) and 655 days (M39/4 366, Labrador Sea) (Fig. 12). The median residence time is 4 days (1.8 days using average weights of encrusted and empty; 3.1 days using average weights of non-encrusted and cytoplasm-bearing average weights for the calculation of shell flux, in which sinking velocity based on shell mass is incorporated). The 95 % confidence interval ranges from 3 days to 5.1 days (1.2 days to 2 days (encrusted and empty); 2.2 days to 3.5 days (non-encrusted and cytoplasm-bearing)), with a geometric mean of 3.9 days (1.5 days (encrusted and empty) 2.8 days (non-encrusted and cytoplasm-bearing)).

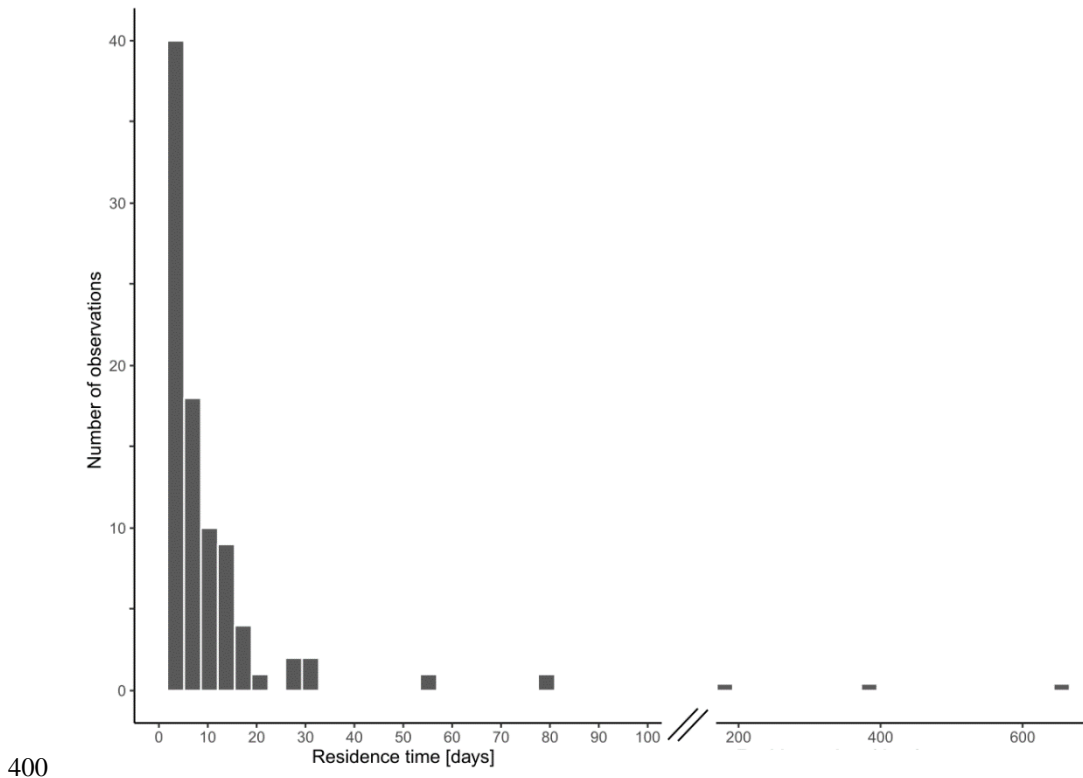



Figure 12: Residence Time of *N. pachyderma* within the productive zone in days, calculated based on the standing stock within the productive zone and the shell flux below the base of the productive zone, calculated with average shall masses below the productive zone. Consider the break in the x-axis between 100 and 200 days.

4 Discussion

4.1 Productive zone and export flux zone

Our analysis of observations from plankton net samples indicates that the productive zone of *N. pachyderma* in the Arctic and Subarctic realm reaches down to about 113 m water depth (median of all samples, 125 m for where calculation after Lončarić et al. (2006) was possible, 136 m defining it as the range end). Greco et al. (2019) have shown that the habitat depth of *N. pachyderma* varies substantially. A variation in the depth interval of maximum abundances of *N. pachyderma* is also presented by Carstens & Wefer (1992) and Carstens et al. (1997), where a connection between distinct water masses and temperature regimes is drawn. Our dataset corroborates these observations and indicates that the base of the productive zone of *N. pachyderma* is also highly variable and reflects the habitat depth (vertical distribution of living specimens). Like Greco et al. (2019), we observe that even if there would be a general pattern of habitat depth and BPZ position being driven by environmental factors, as also proposed by Carstens et al. (1997), it is overlain by considerable variability, even among profiles collected in the same region and around the same time. This means that the observed BPZ variability cannot be driven by the water-column structure alone.

Some of the variability in the BPZ estimates may reflect patchiness in the distribution of planktonic foraminifera populations (Siccha et al., 2012). Meilland et al. (2019) observed that a patchy distribution is mainly present on a horizontal scale, with vertical distribution remaining rather stable. Nonetheless, a horizontally patchy distribution could affect the calculated BPZ in samples from the same region: In profiles with very low shell abundances (< 10, sometimes even < 1 ind. m⁻³), the estimate of the BPZ position may be affected by non-representative estimates of population density. Thus, large abundance differences, caused by a patchy distribution, which has been reported to be best developed for species occurring with high abundances in the Arctic (Meilland et al., 2020), could cause large differences in estimated BPZ and display a variability in the results which ~~might not be as large in reality~~.

In addition, some of the observed variability in BPZ position could arise from differences in sampling methods. Differences in vertical resolution of the compiled plankton net profiles, especially across the BPZ, result in variability that is mainly methodological and which will be reflected in higher variability even among stations from the same region. The BPZ estimate is also affected by the shape of the pattern of change of shell abundance with depth. Where the transition between the productive and the export zone is too gradual, the estimated depth of the BPZ is associated with larger uncertainty.

Some profiles show a pattern of an apparent gain in foraminifera mass flux below the inferred BPZ (Fig. 9). Our analysis of PS93.1 samples indicates that both higher shell abundances and shell weight below the productive zone are present at some of the stations. Higher shell weight could be explained by the loss of lighter, thinner shells due to dissolution, leading to a higher bulk weight at further depth. Gains in fluxes due to higher shell number concentrations are poorly constrained at depths below

the BPZ, as the number of shells present in deeper nets is very low (Fig. A1a-e). A high percentage gain in flux might in some cases only represent a difference of a few shells, which is not related to an actual higher flux but to methodological uncertainties
440 of sampling.

In summary, the calculated BPZ in each profile is associated with some uncertainty. However, we assume that the uncertainties concern all data in the same way, they should not be systematically affecting individual profiles. Therefore, we conclude that the overall pattern of a variable BPZ often located at and even below 100 m, but never below 300 m, should be reliable. These
445 observations helped us to calculate fluxes of shells of *N. pachyderma* from plankton net samples at a more realistic depth in the Arctic Ocean, even where the productive zone has not been explicitly constrained. It can serve as a base for further studies.4.2 Calcification depth

While empty shells are already present in the sampling intervals close to the surface, and the relative abundance of empty shells tends to increase with increasing depth in the productive zone (Fig. 3), shell size does not systematically change with
450 depth (Fig. 5). These observations speak against the presence of extensive OVM by *N. pachyderma* in the studied area (Fig. 2). This is consistent with observations of stable shell sizes of the species with increasing depth in the Barents Sea presented by Ofstad et al. (2020). In contrast, Stangeew (2001) and Manno and Pavlov (2014) described higher abundances of small size fractions in the upper water column close to the surface in *N. pachyderma* from the Fram Strait. However, even in those two studies, some large shells were present in surface samples. Plankton net data from the Nansen Basin from Carstens and Wefer
455 (1992) show higher abundances of small size classes below 100 m depth, which the authors linked to the impact of different water masses in the area. Thus, different conditions at different water depths and/or within different water masses can influence both the abundances of planktonic foraminifera (Carstens et al., 1997) and their size distribution, which could lead to size differences at different depths. The lack of any pattern in shell size in our data does not provide an indication of OVM, and trends in size visible in other studies could in fact be driven by distinct water conditions, and not or not alone by the
460 performance of OVM. Our data also does not present a strong systematic change in size with lunar day, as it was detected in previous studies (Schiebel et al., 2017). However, our shell size data do not cover the entire lunar cycle, preventing drawing firm conclusions on the influence of the lunar cycle on the shell size of *N. pachyderma*.

The likely important role of local environmental parameters on the terminal shell size is also reflected in the differences in
465 shell size between empty and cytoplasm-bearing shells. Empty shells should be representative of specimens that have completed their life cycle. Therefore, shell growth at a constant depth throughout the life cycle of an individual should result in on average larger empty than cytoplasm-bearing shells at all depths. However, we only find such a difference in one of fourteen samples, and on the contrary, significantly bigger cytoplasm-bearing shells occurred in eight of fourteen samples. On the other hand, the calcification intensity of empty shells is significantly higher than for shells bearing cytoplasm in all but one
470 sample, and their shape is significantly more rounded, further indicating strong calcification. This shows that at least in the case of the studied *N. pachyderma*, shell size measured as the maximum diameter of the shell is not an ideal indicator for

maturity, but a highly variable parameter among individual specimens that might reflect variation in environmental conditions during the life cycle of the individual foraminifera. In contrast, the consistently observed stronger calcification intensity of empty shells at all depths and their distinct shape rules out that empty shells in the upper water column only represent specimens

475 affected by premature death (foraminifera which died at a juvenile or young adult stage, likely without reproduction). The stronger calcification compared to cytoplasm-bearing shells is a clear indicator for a completed life cycle, as this species is known to often be associated with the development of a thick terminal calcite layer or crust (Bé, 1960; Kohfeld et al., 1996). In Stangeew (2001), where the presence of OVM is concluded based on shell sizes, the area of occurrence of strongly crusted shells was observed to range from surface to 300 m depth, suggesting reproduction occurred across this whole depth range and

480 not only at its end

Ten out of 18 of the here studied profiles indicate an increase in calcification intensity with increasing depth within the productive zone (Fig. 7), which would speak in favour of OVM. However, with the other half of the profiles not displaying any trend with depth, we must conclude that there is no clear signal for OVM being present or absent. The occurrence of

485 heavily calcified empty shells at all depths indicates that many specimens of *N. pachyderma* reach the final stage in their life cycle, building their final thicker crust, at all depths within their depth habitat. The same conclusion was also favoured by Kohfeld et al. (1996). These authors in addition hypothesised that the local conditions at a given depth not only affect the final size but also calcification intensity. Indeed, like size, calcification intensity in planktonic foraminifera has been shown to reflect parameters like temperature, productivity and optimum growth conditions (e.g. Weinkauf et al., 2016). Those

490 parameters could also cause trends in the calcification intensity with depth, without necessarily being driven by strict OVM. The sampling period of our data has to be considered when evaluating changes of size and calcification intensity with depth: Depending on the life span of *N. pachyderma*, which could be longer than one or two months (Carstens and Wefer, 1992; Kohfeld et al., 1996), it is possible that the samples contain individuals from multiple generations that were produced during different environmental conditions. Furthermore, sinking shells of *N. pachyderma* can be transported over considerable

495 distances, as e.g. shown by v. Gyldenfeldt et al. (2000), whose results would indicate a transport of 25-50 km in the upper 1000 m, resulting in the possibility of some of the encountered specimens being advected from areas with a different hydrography. As this can have an impact on shell size and calcification intensity (e.g. Weinkauf et al., 2016), it could blur signs of OVM. Nevertheless, the estimated residence time of about 4 days in our data indicated that the life span of the sampled *N. pachyderma* is either too short to be strongly affected by environmental condition changes, or that the population size is

500 constant at least across a short time scale. The latter would make huge changes in the environmental conditions unlikely. When interpreting the lower end of estimated residence time, it has to be considered that this might not represent the overall lifespan of the foraminifera, but the days it stays alive after having reached maturity, which is what, based on optical parameters, we mainly analyse. Based on results from culture experiments on *N. pachyderma* from the Southern Ocean, a lifespan of several months, the upper end of our estimated residence time, seems to be possible (Spindler, 1996). It is equally possible that

505 unprecise flux estimations in some samples lead to those extreme values. With the majority of all samples showing a residence

time of only a couple of days, we conclude that the possible blurring of signs of OVM would be rather small, and the lack of a clear trend indicating OVM at all stations can be seen as a reliable result.

If OVM would be present across all specimens of *N. pachyderma* in the Arctic and Subarctic realm, it would need to be very limited in the depth range to not be clearly visible in our data. In regions where the productive zone only starts at about 50 m
510 depth and already ends at about 120 m, the resolution of the studied vertical profiles might be not sufficient to detect it.

In summary, our data on the presence of OVM are inconclusive. The occurrence of empty shells as well as those with high calcification intensity at all depths of the productive zone indicates that *N. pachyderma* does not seem to change its depth habitat during life, while increasing shell calcification could indicate the performance of OVM. Since it seems unlikely that
515 the entire population of the species participates in OVM, we speculate that only a small portion of the specimens follows this behaviour. Indeed, Meilland et al. (2021) suggested such performance of OVM only by a fraction of all specimens within a population for several tropical species in the central Atlantic, and our data would appear to indicate a similar mode of population dynamics for the Arctic *N. pachyderma*. Although our data can neither confirm nor rule out the performance of OVM in *N. pachyderma* in the research area, we can define the calcification zone as the entire upper 300 m of the water
520 column, based on the estimates of the BPZ and the fact that strongly calcified shells can be found within the whole water column above the BPZ.

4.3 CaCO₃ shell mass flux

Knowing the position and variability of the productive zone of *N. pachyderma* in the Arctic, we use data on shell abundances below the productive zone and average shell weights to estimate the calcite flux of *N. pachyderma* in each profile. Estimates
525 of calcite fluxes based on observations from plankton nets are based on two major components that affect the calculations: (i) The (average) shell weight that is used for calculating calcite mass fluxes from shell fluxes and the sinking speed of the shells, and (ii) the depth for which the export fluxes are calculated.

Shell weight varies strongly between different shell types (non-encrusted vs. encrusted) and regions. The shell weight is also
530 influenced by shell size, and the estimates for samples that lack weight measurements are therefore uncertain. Next to possible regional differences in shell sizes, a further source of uncertainty arises from different mesh sizes across the compiled datasets. The used sampling mesh size creates a bias, as a 63 µm net samples different material than a 100 µm or 125 µm net. This bias would result in higher average shell weights when a coarser mesh size is used for sampling. Since we determined and considered weight measurements on samples from the smallest (63 µm) and coarsest (150 µm) mesh sizes, the average that is
535 used in this study is likely representative for most of the samples in the analysed dataset, where most of the profiles are based on sampling with mesh sizes of 100 to 125 µm.

Using weights of encrusted, empty shells results in calcite fluxes that are three to five times higher (average of 10.7 mg CaCO₃ m^{-2} d^{-1}) than estimates based on overall average weights (average of 4.4 mg CaCO₃ m^{-2} d^{-1}) or weights of non-encrusted,

cytoplasm-bearing shells (average of 2.4 mg CaCO$_3$ m^{-2} d^{-1}). However, our observations indicate that not all specimens build a thick crust before reproducing and dying and some still contain remainders of cytoplasm while already sinking. Therefore, flux calculations based on averages of all shell types should be more realistic then only using weights of encrusted, empty shells.

The highest estimated calcite fluxes in our data set are present in the Baffin Bay (average of 13.7 mg CaCO$_3$ m^{-2} d^{-1}; Table 4) and the Greenland Sea (average of 8.0 mg CaCO$_3$ m^{-2} d^{-1}; Table 4). We do not have any weight measurements from the Baffin Bay, hence use overall averages to calculate fluxes for data from there, as explained in the method section (Sect. 2.3). Our data on shell sizes from the Baffin Bay indicate that the shells are systematically smaller than those from the Fram Strait, from where a large number of samples on which weights were measured are taken from. Therefore, the calculated calcite fluxes in the Baffin Bay could be overestimated. As we also see variability in shell weights in samples of similar sizes when sampled at different regions, it would also be hard to establish any size-weight relationship that would be accurate for a region where we lack data. Greenland Sea samples are based on the average weights of samples from the same region, but weight measurements are done on a larger minimum shell size (150 µm) than other counts from the region (mainly using 100 µm mesh size), which could also cause some overestimation.

Interestingly, large differences in flux estimates also emerge from calculations on different depth intervals. We show that flux estimates based on population density at 100 m, as done in previous estimates (e.g. Schiebel, 2002), are overestimating the export, because a large part of the population at that depth is still alive. Calcite mass fluxes based on shell number concentration immediately below the BPZ indicate values that are about five times smaller (8.0 mg CaCO$_3$ m^{-2} d^{-1} below BPZ in contrast to 40.8 mg CaCO$_3$ m^{-2} d^{-1} at 100 m depth; Table 4), indicating that the commonly used level of 100 m (Schiebel 2002) would not be appropriate for the whole Arctic. Next, we observe that the export flux is attenuated below the BPZ (Fig. 9) and average mass fluxes at the deepest sampled net are reduced by a half compared to fluxes directly below the BPZ (average of 4.4 mg CaCO$_3$ m^{-2} d^{-1} at the deepest net). The vertical distribution and amount of this calcite flux loss are similar to observations in other parts of the ocean (Schiebel et al., 2007; Sulpis et al., 2021). Thus, our estimated BPZ seems to be consistent. Sulpis et al. (2021) and Schiebel et al. (2007) ascribe high losses in CaCO$_3$ in the upper water column to indiscriminate digestion by large plankton feeders or CO$_2$ release due to degradation of residual cytoplasm in the shells or in particles to which empty shells may be attached during sinking. Indeed, Greco et al. (2021a) hypothesised that *N. pachyderma* is during life associated with sinking aggregates, which would lead to a situation where even after the foraminiferal cytoplasm is released during reproduction, the empty shell may remain in contact with organic matter. Our data indicate that flux attenuation is driven by both a reduction in shell mass and in shell number concentration (Fig. 9a). Dissolution can result in both of these losses, as both a reduction in weight of strongly calcified shells and a total dissolution of thinner shells is possible due to this process. Notwithstanding the exact mechanisms, our results indicate a substantial attenuation of calcite flux of Arctic *N. pachyderma* below the productive zone, with an average loss of about 6.6 %/100 m. In contrast to other regions, the strong limitation of fluxes in the Arctic to the summer period has to be considered (Bauerfeind et al, 2009; Jonkers et al., 2010). It has been shown

that pulsed high fluxes are less prone to dissolution in the upper water column (Klaas and Archer, 2002; Schiebel, 2002; Sulpis et al., 2021). Therefore, the loss of planktonic foraminifera $CaCO_3$ in the upper water column of the Arctic ocean might be lower than in regions with the same mean annual flux distributed throughout the year.

Based on a compilation of plankton tow data and taking 100 m as the BPZ, Schiebel (2002) reported total planktonic foraminifera calcite flux estimates in the North Atlantic of about 100 mg $CaCO_3$ m^{-2} d^{-1}. This value is more than three times higher than the average calcite export flux by *N. pachyderma* in our dataset at that depth (29.5 mg $CaCO_3$ m^{-2} d^{-1}, averaging over regional averages to account for all regions equally). The difference could be explained by foraminifera building a thicker shell in the North Atlantic, or simply by higher shell abundances. Lower shell abundances in our data already result from methodological effects: By sampling *N. pachyderma* only, we underestimate the total flux of planktonic foraminifera in all regions where abundances of other species are also relevant, like the Greenland Sea and the Norwegian Sea (Fig. 11b). Besides, coarser mesh sizes can underestimate shell number concentrations and hence lead to lower flux values. A comparison of abundances of *N. pachyderma* in our compilation derived from the same region, but sampled with different mesh sizes, shows that its abundance is on average 27 % lower when a coarser mesh size (100 µm, 125 µm, 150 µm) is used, because small shells are not sampled. These observed estimates of a reduction in the abundances is comparable to the results by Carstens et al. (1997), who detected a reduction in foraminifera abundances of 7 % to 40 % with increasing mesh size. The flux given by Schiebel (2002) is based on data from sampling with a 100 µm mesh size. Our data from the western Fram Strait indicates that in this region, the abundance of larger (>125 µm, >150 µm) shells is on average 56 % lower than what is sampled with a mesh size of 100 µm. With 49 out of 148 stations in our dataset having a mesh size coarser than 100 µm, the lower flux estimates in our compilation are likely at least partly underestimated, compared to fluxes consistently based on sampling with a mesh size of 100 µm, but the difference is unlikely to be larger than one third.

Besides, different BPZ at the distinct research areas could lead to different values at 100 m depth. We show that 100 m can be too shallow to estimate the fluxes in the Arctic, but cannot judge the effect of a possibly deeper or varying productive zone in the North Atlantic (Schiebel et al., 1995) on flux estimates. Taking all possible biases in our flux estimation as well as effects on the flux from Schiebel (2002) into account, our estimates cannot be considered as substantially deviating from his flux estimates for the North Atlantic.

An opportunity to further validate our calcite flux estimates is given by a recent study from the Northern Svalbard margin by Anglada-Ortiz et al. (2021), who reported total foraminifera calcite fluxes of 2.3 mg $CaCO_3$ m^{-2} d^{-1} to 7.9 mg $CaCO_3$ m^{-2} d^{-1} based on data from living planktonic foraminifera in the upper 100 m of the water column. It has to be considered that this might not represent the export flux zone, as at least two of the studied profiles show increasing shell abundance below 100 m. Nevertheless, considering that the planktonic foraminifera assemblages reported by those authors contained only about 50% of *N. pachyderma,* their minimum reported flux is similar to the range of the estimates in our data set for the Barents Sea at 100 m (0.39 mg $CaCO_3$ m^{-2} d^{-1} to 1.86 mg $CaCO_3$ m^{-2} d^{-1} using different weight averages for the calculation). That our estimates

are still slightly below those from Anglada-Ortiz et al. (2021), taking the abundance of *N. pachyderma* into account, could be explained by the different mesh sizes: Anglada-Ortiz et al. (2021) sampled with a mesh size of 90 µm, while sampling was done with a mesh size of 125 µm in our data from that region (Table 1). Moreover, the samples analysed in our study were

610 taken in June, while those from Anglada-Ortiz et al. (2021) represent fluxes in August, which often represents the most productive period of planktonic foraminifera in the Arctic Ocean (Jensen, 1998). Overall, this comparison confirms the high local and seasonal variability in fluxes of *N. pachyderma* in the (Sub)Arctic realm (Fig. 11a) and suggests that the estimated flux values in our study are broadly in line with earlier individual observations.

615 To set the estimated flux of *N. pachyderma* into relation to total CaCO$_3$ fluxes of both aragonite and calcite, we compare our results with data from sediment traps in the Greenland Basin (von Bodungen et al., 1995), the Fram Strait (Hebbeln, 2000; Bauerfeind et al., 2009) and the Lomonosov Ridge (Fahl and Nöthig, 2007). As all of our data originate from the summer season, and the shell flux in the Arctic and Subarctic is highly seasonal (Jensen, 1998), we compare our data with daily CaCO$_3$ fluxes from June to September only. The total range of CaCO$_3$ fluxes is similar to the flux we observe in *N. pachyderma* in

620 plankton nets, with fluxes of *N. pachyderma* being mostly located at the lower end (Fig 13). Using a mean daily mass flux of *N. pachyderma* at the greatest sampled depths of each net of 4.43 mg CaCO$_3$ m^{-2}, the species would make up about 23 % of total CaCO$_3$ flux (18.89 mg CaCO$_3$ m^{-2}) measured in the sediment traps. This is in line with global estimates from Schiebel et al. (2007) giving a contribution of planktonic foraminifera to overall CaCO$_3$ fluxes of about 25 %. Our result is further in line with an estimated contribution of planktonic foraminifera to total CaCO$_3$ fluxes in the Atlantic Ocean of 19 % by Kiss et al.

625 (2021), but lower than estimates from Salmon et al. (2015) of up to 40 %. For the Southern Ocean, higher contributions (34-49 %) have been estimated (Salter et al., 2014).

A direct comparison of fluxes of planktonic foraminifera from samples from within the same region with total CaCO$_3$ fluxes in the region indicates a lower contribution of planktonic foraminifera in the Eastern (> 0°) Fram Strait (10 %) and a higher contribution in the western part (< 0° E) of the Greenland Sea (50 %) to total CaCO$_3$ fluxes. For this comparison, we subdivided

630 the regions by longitude to account for the different influences of Atlantic and Arctic waters, which play an important role for the abundances and habitats of planktonic foraminifera in this region (Pados & Spielhagen, 2014). The contribution of 10 % in the Fram Strait is in line with the lower end of estimated contribution of planktonic foraminifera to total CaCO$_3$ fluxes at the Northern Svalbard margin (4-34 %; Anglada-Ortiz et al., 2021). The higher contribution in the Greenland Sea is in the range of the estimates from Salter et al. (2014) from the Crozet Plateau in the Southern Indian Ocean, indicating that the given

635 contribution falls within globally realistic ranges. The previously described possible effect of coarser mesh size decreasing flux estimates has to be considered, meaning that the values from our dataset provide a minimum range.

Overall, our data indicates that the production of CaCO$_3$ by planktonic foraminifera in the Arctic Ocean has a similar share to total fluxes as in other regions. We also see large variability with some Arctic regions showing much lower contribution than in other oceans and the global average. It has to be stressed, however, that our estimates are only for a single (albeit often the

640 most abundant) species and the total flux of planktonic foraminifera in the studied region must be higher. The contribution of

planktonic foraminifera to the Arctic carbonate budget may therefore be larger than the numbers given here. Moreover, even though the aragonite-producing pteropods are abundant in the Arctic and their shells are preserved in sediment trap samples (Bauerfeind et al., 2014; Busch et al., 2015), most of the aragonite flux dissolves prior to burial in the sediment because the majority of the Arctic seafloor is located below the aragonite compensation depth (Jutterström and Anderson, 2005). Our

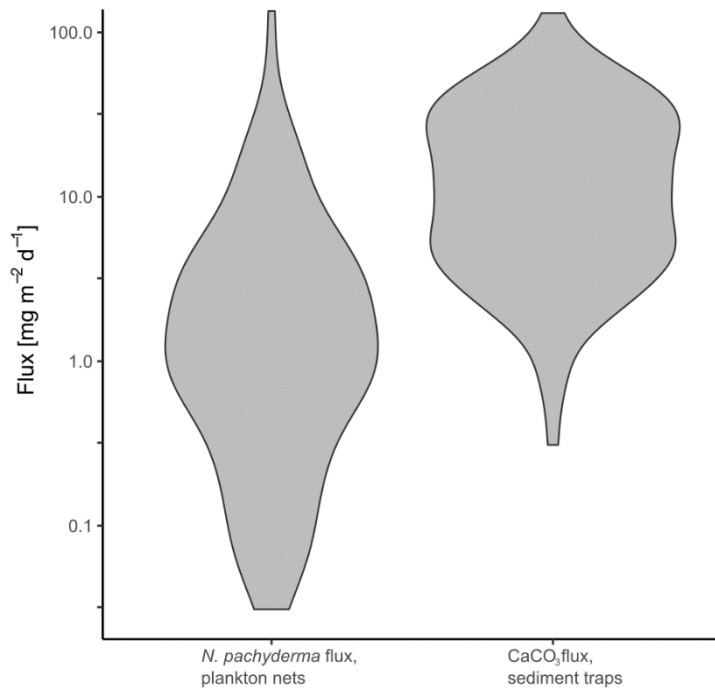
645 calculation of an apparent 23 % contribution of planktonic foraminifera to the summertime export flux of carbonate is thus likely translated into a larger share of the burial flux, making the calcite flux by planktonic foraminifera highly relevant for the Arctic oceanic carbon cycle.

650

Figure 13: Comparison of logarithmic daily mass flux of *N. pachyderma* in plankton nets and CaCO₃ in sediment traps. Sediment trap data is from sediment traps in the Fram Strait (Hebbeln, 2000; Bauerfeind et al., 2009), the Greenland Basin (von Bodungen et al., 1995) and the Lomonosov Ridge (Fahl and Nöthig, 2007).

5 Conclusion

655 Our compilation of vertically resolved data on the dominant Arctic planktonic foraminifera *N. pachyderma* reveals that the base of the productive zone of this species is on median located at about 113 m depth, but shows large regional variability and locally reaches down to 300 m. Our analyses show that it is important to constrain the base of the productive zone to estimate

fluxes in the export flux below: using a constant 100 m depth to estimate fluxes leads to a fivefold flux overestimation in contrast to the flux at the top of the export zone. Below the BPZ, the shell flux is decreasing on average by 6.6 % per 100 m,

660 with highest losses directly below the BPZ. No further change is observed deeper than 300 m below the BPZ. Therefore, we can conclude that in the absence of knowledge on the position of the BPZ, using 300 m depth should provide a conservative, yet more realistic estimate of the *N. pachyderma* export flux in the Arctic realm than using the formerly often used depth of 100 m. Within the productive zone, our data are inconclusive whether *N. pachyderma* performs ontogenetic vertical migration throughout its lifecycle. We observe empty and strongly encrusted shells, hence specimens that have completed their life cycle,

665 at the whole depth range, and do not see any pattern of increasing shell size. Nevertheless, as a systematic increase in calcification intensity with depth is present at some stations, we speculate that OVM is performed by at least a small part of the community.

The overall average calcite mass flux of *N. pachyderma* based on measured average shell weights (average of 3.4 µg) and shell number concentrations (average of 25 ind. m^{-3}) is estimated to be 8 mg CaCO$_3$ m^{-2} d^{-1} directly below the base of the productive

670 zone in the Subarctic and Arctic Ocean. From this depth on, the flux on average is attenuated at a rate of 6.6 %/100 m at least within the following 300 m depth. This attenuation is driven by a reduction in shell number concentration and weight, which is probably mainly driven by dissolution of thinner, less calcified shells.

Notwithstanding uncertainties in flux estimates due to high regional variability, coarser mesh sizes with underrepresentation of total shell abundance and the lack of weight measurements in some regions, our estimates are in line with previous global

675 studies and local studies from adjacent areas. Comparison with data from sediment traps shows that *N. pachyderma* is on average responsible for 23 % of total pelagic carbonate fluxes in the Subarctic and Arctic realm, with a regional variability of ten to 50 %, indicating an even bigger share of total planktonic foraminifera especially in Subarctic regions, where *N. pachyderma* only makes up 50 % of the total population.

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Data availability

All data created for this study, including calculations made on published data, is available on PANGEA (https://doi.pangaea.de/10.1594/PANGAEA.941250). The data source of abundances from published data are also listed in Table 1, and shell weights from the M39/4 expedition are listed as PANGEA references.

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Author contributions

The study was designed by all authors. FT carried out the laboratory work with help from JM and the data analysis with help from LJ. All authors contributed to the interpretation and discussion of the results. FT wrote the paper with contributions from MK, LJ and JM.

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Competing interests

The authors declare that they have no conflict of interest.

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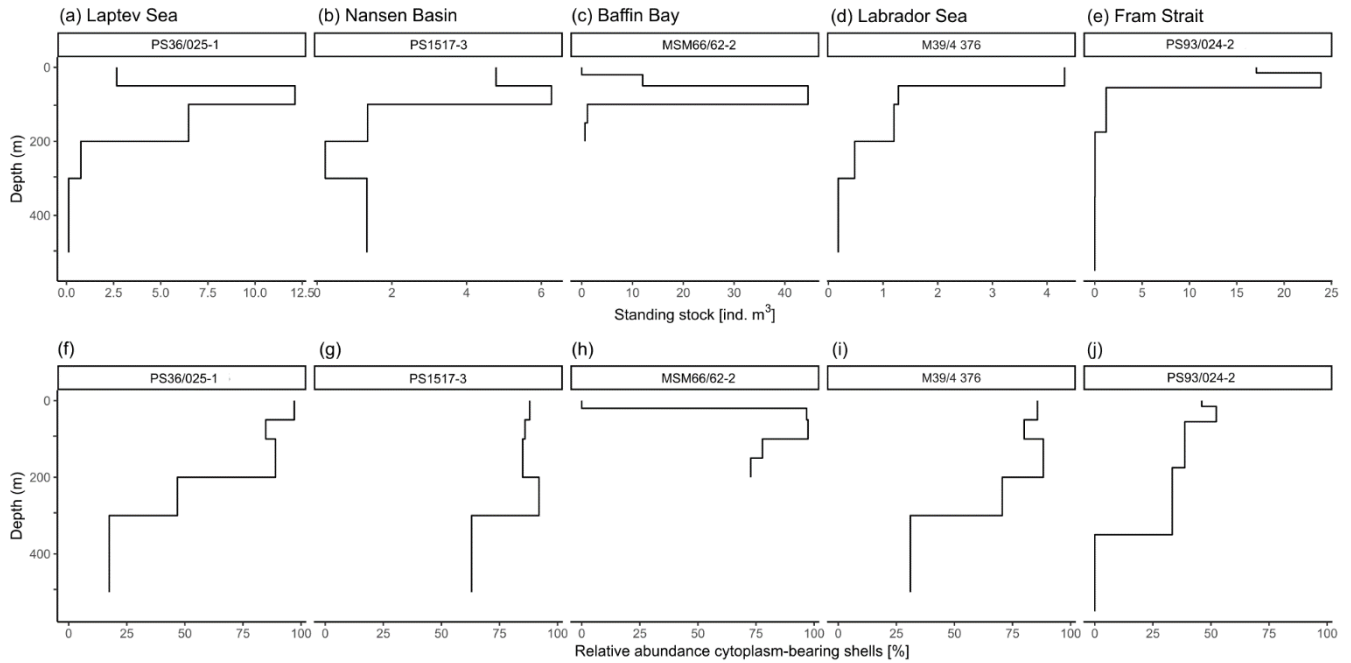
Appendix A

900

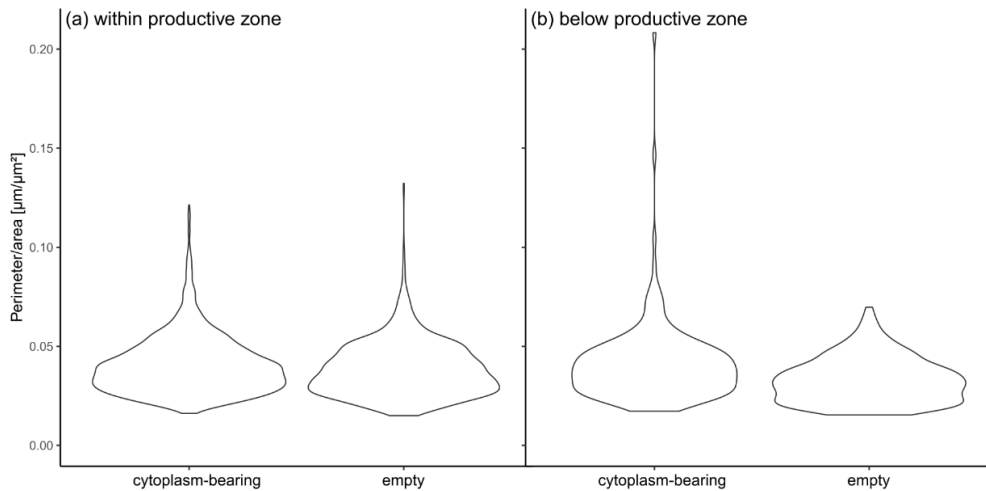
Figure A1: Example of vertical profiles of abundances of *N. pachyderma* at five different sampling locations from different parts of the Arctic Ocean. (a-e) show absolute shell number concentration (ind. m^{-3}) of *N. pachyderma*, (f-j) relative abundance of cytoplasm-bearing shells of *N. pachyderma*.

905

Figure A2: The ratio of perimeter to area in individual shells of *N. pachyderma* in samples from PS 93.1, divided by the status (cytoplasm-bearing and empty). (a) represents the shells from within the calculated productive zone of the individual stations, (b) those from below the productive zone.

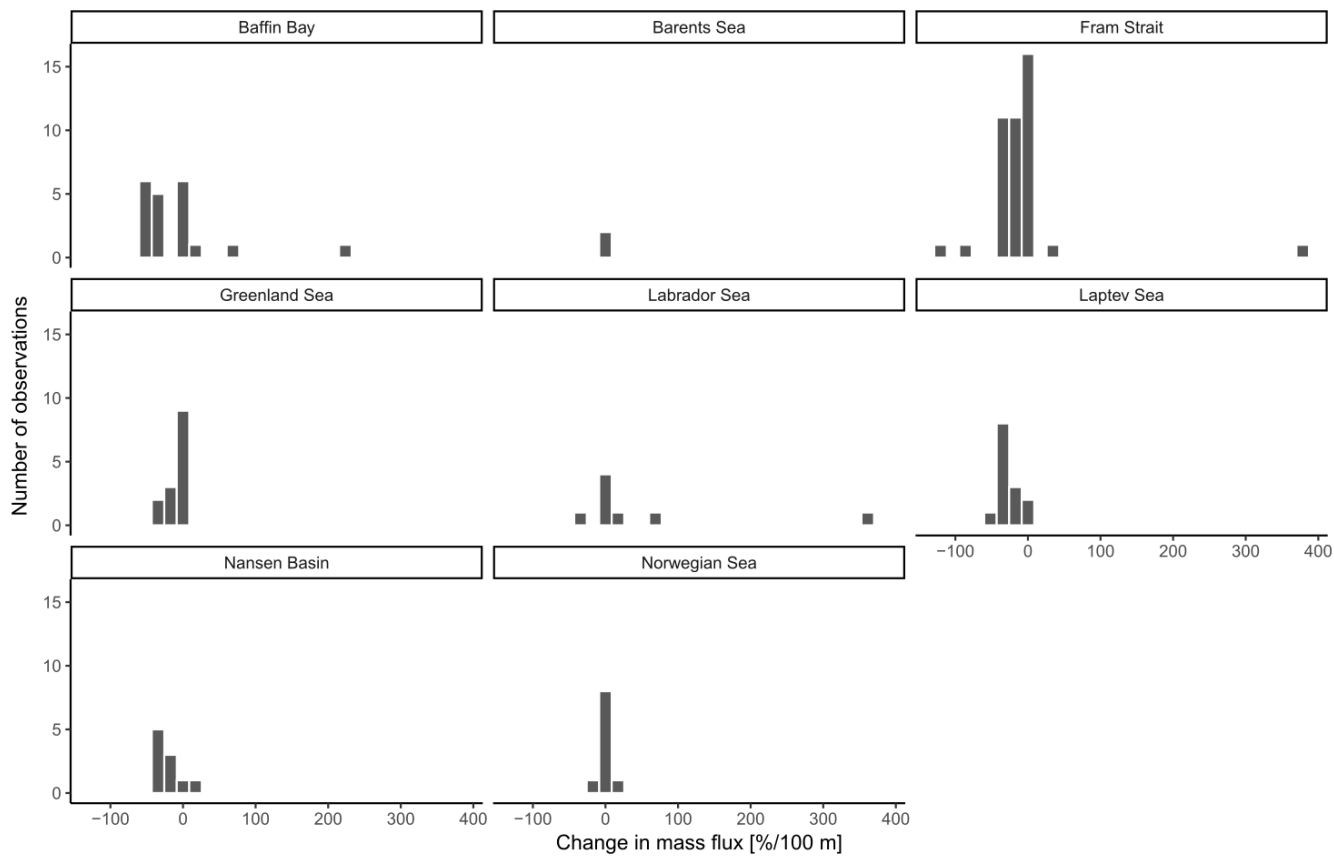



Figure A3: Loss in shell flux between the net directly below the calculated base of the productive zone and the deepest net of sampling of each station, divided by the different regions of sampling.