



Benthic Alkalinity fluxes from coastal sediments of the Baltic and North Seas: Comparing approaches and identifying knowledge gaps

Bryce Van Dam^{1,*}, Nele Lehmann^{1,4,7} Mary A. Zeller³, Andreas Neumann¹, Daniel Pröfrock², Marko Lipka³, Helmuth Thomas^{1,7}, Michael E. Böttcher^{3,5,6}

- ¹ Helmholtz-Zentrum Hereon, Institute of Carbon Cycles, Geesthacht, Germany
- ² Helmholtz-Zentrum Hereon, Institute of Coastal Environmental Chemistry, Geesthacht, Germany
- ³ Leibniz Institute for Baltic Sea Research (IOW), Warnemünde, Germany
- ⁴Alfred Wegener Institute Helmholtz Centre for Polar and Marine Research, Potsdam, Germany
- 10 5 Marine Geochemistry, University of Greifswald, FRG
 - ⁶ Interdisciplinary Faculty, University of Rostock, FRG
 - ⁷ University of Oldenburg, Oldenburg, Germany

Abstract. Benthic alkalinity production is often suggested as a major driver of net carbon sequestration in continental shelf ecosystems. However, information and direct measurements of benthic alkalinity fluxes are limited and are especially challenging when biological and dynamic physical forcing causes surficial sediments to be vigorously irrigated. To address this shortcoming, we quantified net sediment-water exchange of alkalinity using a suite of complementary methods, including 1) 224 Ra budgeting, 2) incubations with 224 Ra and Bromide as tracers, and 3) numerical modelling of porewater profiles. We choose a set of sites in the shallow southern North Sea and western Baltic Sea, allowing us to incorporate frequently occurring sediment classes ranging from coarse sands to muds, and sediment-water interfaces ranging from biologically irrigated and advective to diffusive into the investigations. Sediment-water irrigation rates in the southern North Sea were approximately twice as high as previously estimated for the region, in part due to measured porewater 224 Ra activities higher than previously assumed. Net alkalinity fluxes in the Baltic Sea were relatively low, ranging from an uptake of -35 μ mol m⁻² hr⁻¹ to a release of 53 μ mol m⁻² hr⁻¹, and in the North Sea from 1 to 33.6 μ mol m⁻² hr⁻¹. Lower than expected apparent nitrate consumption (potential denitrification), across all sites, is one explanation for our small measured net alkalinity fluxes. Carbonate mineral precipitation and sulfide re-oxidation also appear to play important roles shaping net sediment-water fluxes in the North Sea and Baltic Sea sites, respectively.

1 Introduction

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Continental shelf systems are considered to act as sinks for atmospheric CO₂ (Borges et al., 2005; Laruelle et al 2010; 2018; Rutherford et al., 2021), with an important but currently uncertain role in the global carbon budget (Lacroix et al., 2021). Current trends in shelf CO₂ uptake are highly variable (Laruelle et al., 2018), in response to a variety of factors, including: regional trajectories in primary production, acid-base buffering (Thomas et al., 2007), the underlying geology, mixing with

^{*}Correspondence to: Bryce Van Dam (Bryce.Dam@hereon.de)



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riverine and open-ocean waters, and wind patterns (Meyer et al., 2018). Where water depths are relatively shallow, there is an increased potential for benthic-pelagic coupling to affect the net shelf CO₂ uptake. In regions like the shallow southern North Sea, persistent wind and tidal-driven mixing strongly impacts both the benthic and pelagic ecosystems. This mixing ventilates surface sediments, supplying oxidants for the respiration of organic matter and flushing out the reduced products, thereby affecting net sediment-water exchange of dissolved substances like inorganic carbon (DIC) and total alkalinity (TA). Aerobic remineralization is a key DIC source in physically or biologically ventilated surface sediments (Neumann et al., 2021; Rassmann et al., 2020), followed by microbial sulfate reduction (MSR; Al-Raei et al., 2009; Werner et al., 1999, 2001). These DIC sources may be further enhanced to varying extents by anaerobic processes like denitrification, as well as iron and manganese reduction, which, like MSR also produce TA in addition to DIC (e.g., Zeebe & Wolf-Galdrow, 2001). While intense biological reworking limits net organic carbon storage in North Sea sediments (Diesing et al., 2020; de Haas et al., 1997), net carbon uptake from the atmosphere may be possible through a combination of the shelf CO₂ pump (Thomas et al., 2004), as facilitated by large oceanic inflows (Lacroix et al., 2021), as well as a long-term increase in the DIC inventory driven by combined biological and anthropogenic CO₂ (Clargo et al., 2015).

Additionally, there is a suite of geochemical processes which may also affect net shelf CO₂ uptake. For example, the precipitation and dissolution of calcium carbonate (CaCO₃) and silicate minerals, pyrite formation and oxidation, dissolution and formation of silicate minerals, and reverse weathering may substantially affect net sediment-water fluxes of DIC and TA, and, hence, the carbonate chemistry of the overlying water column (Berelson et al., 2019; Hagens et al., 2015; Pätsch et al., 2018; Rassmann et al., 2020; Thomas et al., 2009; Winde et al., 2017; Schwichtenberg et al., 2020). While not a direct result of biological metabolism, the kinetics of these abiotic processes are determined by the biologically mediated redox setting impacting the physico-chemical saturation states (Morse and Mackenzie, 1990). Therefore, a strong interplay exists in shelf sediments, between biogeochemical cycling of macro and minor nutrients, and benthic (bio)geochemical solid-solution interactions, all of which is further mediated by wind- and tidal-driven physical mixing.



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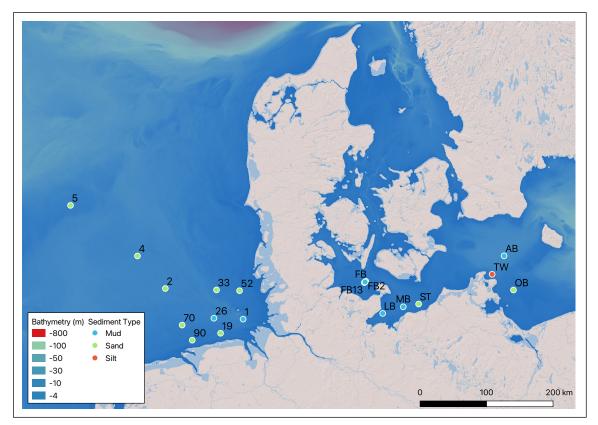


Figure 1. Map of sampled stations, with points colored by sediment type.

Substantial prior work in the coastal North Sea shows that these sediment (bio)geochemical processes exert a strong cumulative effect on water-column carbonate chemistry (Brenner et al., 2016; Burt et al., 2014) and ultimate CO₂ exchange with the atmosphere (Thomas et al., 2009). Recent modelling and experimental efforts point to benthic denitrification of riverine nitrate as a dominant TA source to the southern North Sea, but also imply a substantial role of benthic carbonate mineral dissolution (Winde et al., 2014; Pätsch et al., 2018; Burt et al., 2016; Schwichtenberg et al., 2020). In addition to these processes, metal (Fe and Mn) reduction can also be an important TA source, especially in depositional centers with finergrained sediment and greater organic carbon loads (Lenstra et al., 2019; Reithmaier et al., 2021). However, rapid sediment accumulation may decouple net Fe reduction from sediment-water TA fluxes (Rassmann et al., 2020). Hence, more investigations of the benthic (bio)geochemical processes in relation to benthic DIC/TA fluxes is clearly warranted.

The Baltic and North Seas form a continuum of water and material transport (Gustafsson 1997; Kuliński et al., 2022; Maar et al., 2011), deeply linking the carbon cycles of these two ocean basins. While the nearshore regions of the southern North and western Baltic Seas are located at a similar latitude, have a similar water depth, and contain similar sediment types, they experience very different physical forcing. While tidal forcing dominates in the southern North Sea, the western Baltic is instead characterized by negligible tidal forcing, estuarine mixing and associated salinity variability. For example, large tidal





forcing combined with coarse-grained sediments in the southern North Sea, could promote advective over diffusive fluxes, and increase the oxygen penetration depth (Werner et al., 1999, 2003; Billerbeck et al. 2006; Al-Raei et al., 2009). Bioturbation and bioirrigation are also especially important in the coastal North Sea and can explain how sediment-water fluxes of oxygen and other elements vary across season and sediment type (Lipka et al., 2018; Gogina et al., 2018; Neumann et al., 2021; Bratek et al., 2020). In contrast, limited tidal forcing combined with finer-grained sediments in the western Baltic Sea could promote stronger biogeochemical zonation and a relatively greater importance of anaerobic over aerobic processes (Böttcher et al., 2000). While we know that these factors shape benthic biogeochemical processing, sediment-water fluxes in these shallow coastal environments are still poorly-constrained, limiting our understanding of their role in the regional carbon cycle (Kuliński et al., 2022).

In this study, we present new results quantifying sediment-water DIC and TA fluxes in the southern North Sea, considering a variety of methods, including Ra budgets and core incubations. This dataset is augmented by archived porewater data from the Baltic Sea and some additional sites in the North Sea, in order to further understand the differential impacts of physical and biogeochemical forcing in these two nearby basins. Benthic TA, DIC, and element fluxes were modelled from these porewater profiles, aimed at predicting net sediment-water fluxes of DIC and TA across a broad range in sediment types. The relationships of especially SO₄²⁻, DIC, TA, Fe, and PO₄³⁻ were utilized as a first approach on the potential role of secondary (bio)geochemical processes described above, which are so far only poorly represented in regional carbon budgets (Pätsch et al., 2018; Schwichtenberg et al., 2020), in the context of the coupled physical and biological drivers of net benthic-pelagic coupling. Each applied method has its own advantages and disadvantages to emphasize advection, diffusion, or bioirrigation to varying degrees. The differences between the approaches are partially related to methodological limits, but may also indicate real differences between the consequences of physical and biological drivers on biogeochemical processes such as net CO₂ uptake and, therefore, require further investigations.

2 Methods

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2.1 Dataset description

Samples collected for this study come from a set of 19 stations in the southern North Sea (10 sites) and the western Baltic Sea (9 sites) that were visited during cruises between 2015 and 2020 (Figure 1). Here we present results from four cruises where sampling and experimental approaches are comparable; these were HE541 (Sept 2019) and MSM50a (Jan 2016) in the North Sea, and EMB111 (Aug-Sept 2015) and EMB238 (May 2020) in the Baltic Sea. Parts of the experimental results from MSM50 and EMB111 have previously been published by Lipka et al. (2018) and Gogina et al. (2018), and these sites were chosen to be complementary in terms of water depth, distance from shore, and sediment type. Only results from HE541 in the southern North Sea are used for the Ra budget and core incubations, while porewater profile modelling and constituent ratio analyses are given for all sites. The water depth at these sites is shallow at 23.6 +/- 8.5 m (mean +/- S.D.), with a maximum



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of 46.0 m (Site AB) and minimum of 11.2 m (Site 52). Of the 19 stations considered here, 10 are classified as sandy sediments, 8 as muds, and one as silt sediment (Lipka, 2017; Lipka et al., 2018).

At each site, a multicoring device was used to collect 10 cm diameter cores which contained approximately 20 cm of sediment. Sediment porewaters were extracted within 6 hrs after core collection using Rhizons (nominal pore size of $0.15 \mu m$) at 1-2 cm intervals for the first 10 cm, and with a typical interval of 5 cm below that. Table S1 in the Supporting Information lists the preservation techniques and measurement methods for each analyzed parameter.

2.2 Lab Analyses

The samples taken during the HE541 cruise were analyzed according to the following protocols: TA was determined at IOW via a small volume, two-point potentiometric titration according to van den Berg and Rogers, 1987: The sample (400 μ L) was transferred into a vial with a pre-set volume of acid on board, assuming that all weak acids in solution would be protonated (pH < 3). The sample was cooled at 4 °C until further analysis. Back in the lab, the sample was allowed to equilibrate to room temperature and the first potentiometric point was measured using a pH meter (SevenMulti, Mettler-Toledo, Gießen, Germany). After the addition of HCl (75 μ L, 0.1 M), the second point was recorded (pH \approx 2). Taking these two points, the slope of the electrode response was determined (van den Berg and Rogers, 1987). The measurements were calibrated against certified reference materials (CRMs), from batch 142 provided by Prof Andrew Dickson (Scripps Institution of Oceanography, San Diego, USA).

DIC and δ^{13} C-DIC were also analyzed at IOW with a continuous-flow isotope-ratio-monitoring mass spectrometry (CF-irmMS) using a gas mass spectrometer (Finnigan MAT 253, Thermo Fisher Scientific, Waltham, USA) coupled to a gas bench (GasBench II, Thermo Fisher Scientific, Waltham, USA) via a continuous flow interface (ConFlo IV, Thermo Fisher Scientific, Waltham, USA) (Winde et al., 2014a). A calibration of the instrument was performed against CRMs provided by Prof Andrew Dickson (Scripps Institution of Oceanography, San Diego, USA). Sulfide concentrations were measured at IOW using the methylene blue technique (Cline, 1969) on a Specord 40 a spectrophotometer (Analytik Jena, Germany).

Determination of elemental concentrations (Ca, Fe, Mn, P, S) was performed at Hereon using an ICP-MS/MS (Agilent 8800, Agilent Technologies, Tokyo, Japan) coupled to an ESI SC-4 DX FAST autosampler (Elemental Scientific, Omaha, Nebraska, USA) equipped with a discrete sampling system with a loop volume of 1.5 mL (Zimmermann et al., 2020). Measurements were validated with a seawater standard (S = 35, IAPSO Standard Seawater, Ocean Scientific International Ltd, Hampshire, United Kingdom) with the addition of Fe and Mn (c = 1 mg/L). Calibrations were prepared from either single or multi element solutions traceable to NIST CRMS. Nutrient concentrations were determined using an automated continuous flow system (AA3, Seal Analytical, Norderstedt, Germany) and standard colorimetric techniques (Graßhoff and Almgren, 1983). Dissolved metals, S, P, and Si results from the Baltic Sea samples were measured at IOW via ICP-OES at IOW as described by Winde et al. (2014a) and Lipka et al. (2018).



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2.3 Sediment-water flux determinations

In this study, we applied three independent methods to estimate sediment-water fluxes of major, minor, and trace elements. This is a conservative approach, because each individual method emphasizes advective or diffusive processes to a different degree, ensuring that our calculated sediment-water fluxes span a complete range in possible rates.

First, a water-column ²²⁴Ra decay balance, used to derive porewater irrigation rates, which are in turn used to parameterize bulk sediment-water fluxes based on measured concentrations in the upper porewaters. The second approach is similar to the first, but with irrigation rates derived from ship-board core incubation experiments, with both bromide and ²²⁴Ra employed as tracers. We also used PROFILE models to assess sediment-water fluxes for all sites, which uses the shape of porewater profiles to balance internal production and consumption with a vertical diffusive transport (Berg et al., 1998).

2.4 ²²⁴Ra decay balance

Samples for ²²⁴Ra were collected exclusively during the HE541 cruise in the southern North Sea, following the methods of Moore et al. (2011) and Burt et al., 2016 (²²⁴Ra determinations were not made during the other cruises). Briefly, approximately 100 L of water from the ship's seawater line was passed through a cascade of 10 and 1 µm filters, then pumped slowly (1 L/min) through a cartridge containing manganese oxide-coated fibers which quantitatively adsorb Ra (confirmed with efficiency samples). After rinsing and drying, samples were counted first within 24 hours of collection on board the ship with a Radium Delayed Coincidence Counting (RaDeCC) system (Moore and Arnold 1996). A second count was executed after 7-14 days, allowing us to apply the error propagation technique of Burt et al., 2016 to derive uncertainty statistics for the final ²²⁴Ra activities given in section 3.1. In addition to the bulk seawater samples, porewaters were also collected from all sites during the HE541 cruise using Rhizons, with water from all core depths combined into a single sample (~100-200 mL) that was treated and counted on a RaDeCC as described above.

Because ²²⁴Ra is not a gas and has a relatively short half-life (3.7 days), its activity in surface waters is a relatively simple balance between advective-diffusive inputs from the sediments (where it decays from its longer-lived parent isotope ²²⁸Th), lateral exchange in surface waters, and radioactive decay (Garcia-Orellana et al., 2021). We assume that lateral exchange is a small term in this budget, allowing us to calculate the benthic advective-diffusive input term (net irrigation) as the balance between radioactive decay and the measured surface water activity. These irrigation rates derived from the ²²⁴Ra decay balance (mean = 81 L m⁻² d⁻¹) were then used to calculate net sediment-water fluxes by multiplying by the average concentration difference between the pore water and the overlying water concentration. For "sandy" sites, we applied the average porewater concentration of each parameter from the upper 5 cm in each core, while a single concentration from the top sediment layer of each core was used for the "muddy" and "silty" sites. Sign convention is that positive fluxes indicate a flux out of the sediment and vice versa.



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2.5 Ship-board incubations

Benthic fluxes were measured on intact sediment cores from Multicorer by means of whole core batch incubations. Typically, 3 to 4 intact sediment cores in transparent plastic liners (PMMA, 10 cm inner diameter, 60 cm length) per station were selected, which had no visible perturbations such as cracks, voids, or injured animals. The sediment cores were typically 15 - 30 cm in length. The incubation was enclosed with a gas-tight lid that was adjusted to a resulting supernatant height of 15 cm (approx. 1 L volume). The water column was constantly stirred by horizontal propellers. The stirring intensity was adjusted to the highest intensity that did not result in sediment resuspension to ensure vigorous mixing of the supernatant and prevent suspended particles from settling. Incubations were executed aboard the research vessel in a temperature-controlled lab set to in-situ temperature. Primary production was excluded by wrapping the cores in aluminum foil. At the beginning of each incubation, a NaBr solution was injected into the supernatant, which resulted in a final Br concentration of 1.6 mmol / L. During incubation, oxygen was monitored continuously to assess the progress, and the incubation was terminated when the oxygen saturation dropped below approximately 80% (typically after 12 to 24 h). Water samples for Bromide analysis were drawn from the core supernatant in 3-6 h intervals (typically 10 time steps) by means of syringes connected to PVC-tubing. Water samples were then filtered through 0.45 μ m syringe filters and stored frozen until analysis in the land-based laboratory. After incubations, additional samples form the supernatant were taken for measurements of Radium and DIC. The incubation method is described in detail in Neumann et al. (2021).

2.6 PROFILE Modelling

We first employed the one-dimensional numerical model tool, PROFILE, to derive sediment-water fluxes and internal rates of production and consumption for most measured constituents (Berg et al., 1998). PROFILE builds a best-fit model from measured porewater profiles, assuming they are representative of steady state conditions. This model is then split into several equidistant zones (typically 5 to 10 discrete intervals), each of which is fit with a specific net production rate. We then set a first boundary condition as the calculated diffusive flux out the bottom of the model domain, using the concentration gradient in the two deepest porewater samples, as well as the sediment porosity and calculated the molecular diffusion coefficient (Table 1). The final sediment-water flux is then determined through conservation of mass between net internal production and the bottom diffusive flux. Further parameters needed to inform PROFILE include: sediment porosity (φ) , biodiffusivity, irrigation, diffusivity, statistical terms, and others given in Table 1.

Table 1. Parameterization of PROFILE model

		Sand	Silt	Mud
Max deviation when accepting value (%)		0.001	0.001	0.001
Level of significance for F-statistic		0.2	0.2	0.2
Biodiffusivity (cm ² hr ⁻¹)	< 5 cm	0.36	0.18	0.18
	5-10 cm	0.18	0	0
	>10 cm	0	0	0





Irrigation coefficient (hr ⁻¹)	< 5 cm	0.11	0.0011	0.0011
	5-10 cm	0.011	0	0
	>10 cm	0	0	0
Porosity (φ)		0.4	0.6	0.8
Coefficient of Molecular Diffusivity (cm ² hr ⁻¹)		Calculated in R: 'marelac' as f(Sal, Temp, Pres)		
Sediment Diffusivity (cm ² hr ⁻¹)		$D_s = D/(1 + 3[1 - \varphi])$		
Boundary Conditions		Concentration and diffusive flux at the bottom		

3 Results and Discussion

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3.1 North Sea Irrigation rates (decay balance and incubation)

Surface water 224 Ra activities were consistent throughout the southern North Sea, ranging from 248 ± 11.6 dpm m⁻³ (Site 33) to 384 ± 21.3 dpm m⁻³ (Site 90). This is approximately twice as high as found in previous studies in the region (Burt et al 2014; 2016), but similar to activities for the Wadden Sea (Moore et al., 2011). Our observation of high surface water 224 Ra is consistent with early fall wind-driven vertical mixing causing an enhanced exchange between porewater high in 224 Ra and the surface water. The wind-driven water column mixing also had the effect of breaking down any vertical stratification that may have developed during the summer (not shown). Our porewater 224 Ra activities (8.4 ± 0.89 to 44 ± 3.0 dpm L⁻¹; average of 19.9 dpm L⁻¹ (or 19,946 dpm m⁻³)) were also approximately double compared to previous measurements in the German Bight (Burt et al., 2014), and the Wadden Sea (Moore et al., 2011). These high 224 Ra activities yield calculated irrigation rates (mean = 81 L m^{-2} d⁻¹; red bars in Figure 2) that were about twice the levels found by Burt et al., (2014). Solute fluxes for these sites are derived using PROFILE modelling, as discussed later. The high variability in porewater 224 Ra activity (8.4 to 8.4 dpm L⁻¹) also shows that care should be taken when deciding on endmember activities for 224 Ra-based budgets aimed at assessing porewater irrigation rates (Cook et al., 2018; Garcia-Orellana et al., 2021).

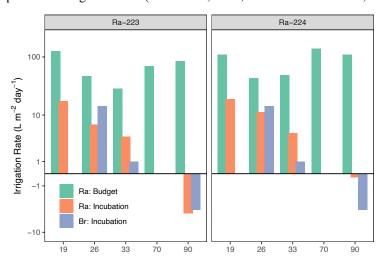






Figure 2. Porewater irrigation rates (L m⁻² d⁻¹) derived from Ra budget (green) and incubations with Ra (orange) or Bromide (blue) as tracers.

Irrigation rates derived from Br and ²²⁴Ra measurements during the ship-board incubation experiments (mean = 10.5 L m⁻² d⁻¹; blue bars) were a factor of 8 lower than rates derived from the ²²⁴Ra decay balance. This is consistent with prior studies in the southern North Sea, where incubation-based fluxes were 2-3 times lower than decay-balance estimates (Burt et al 2014). This difference is likely due to the fact that the decay balance implicitly represents the combined diffusive, advective, and bio-irrigative processes that exist in the environment, which are incompletely captured in whole-core incubations. We must stress here that the Ra decay-balance does not necessarily represent a better estimate of the real irrigation rate, as many issues also exist with this method especially related to porewater endmember determination and the assumption of steady state (Cook et al., 2018; Garcia-Orellana et al., 2021). Rather, these irrigation rates and solute fluxes presented below, simply represent a most likely range in "real-world" values. The difference among approaches can also illustrate the varying importance of physical and bio-mediated forcing of net sediment-water exchange.

220 3.2 Net TA Fluxes

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3.2.1 North Sea TA fluxes from irrigation, incubation, and PROFILE modelling

Using the range in irrigation rates shown above, we calculated net sediment-water fluxes for a variety of solutes using the measured surface-pore water concentration difference (by convention, positive is out of the sediment). Average TA fluxes are tabulated in table 2. At only two sites in the North Sea was porewater TA enriched enough to reliably calculate net fluxes using the incubation or budget approaches, given the relatively low precision of the small-volume TA titration method applied here. However, at these two sites (19 and 26), our TA fluxes (Figure 3) were 1-2 orders of magnitude lower than previous estimates for this region (Brenner et al., 2016; Burt et al., 2014; Voynova et al., 2019). Excluding the PROFILE-modelled TA fluxes for the North Sea (which had very low R²), net TA fluxes in the North Sea were relatively low, ranging from 1 to 6.3 µmol m² hr¹ across all sites and sediment types (Table 2).

Table 2. Comparison of average TA fluxes (μ mol m⁻² hr⁻¹) between methods in the North and Baltic Seas, separated by sediment type.

Basin	Method	Mud	Sand	Silt
Baltic Sea	PROFILE	53 ± 95.3 (n=6)	$-4.3 \pm 43.5 \text{ (n=2)}$	-35 (n=1)
North Sea	PROFILE	$1 \pm 2.1 (n=3)$	$33.6 \pm 42.3 \ (n=5)$	-
North Sea	Ra Incubation	$1.5 \pm 3.5 \ (n=2)$	1.1 ± 1.5 (n=2)	-
North Sea	Ra Budget	$5.6 \pm 13.4 (n=2)$	$6.3 \pm 8.8 \ (n=2)$	-
North Sea	Br Incubation	1.8 ± 4.4 (n=2)	-	-



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Burt et al., (2014) applied a similar Ra-based approach to estimate benthic fluxes but used a porewater-surface water TA difference of 0.7 mM, far greater than what we observed. For example, our porewater TA measurements at sites 19 and 26 were on average 2.23 and 2.26 mM, only 2 - 4 μM (0.002 – 0.004 mM) greater than bottom water TA. As a result, the inferred sediment-water TA fluxes in Burt et al., 2014 were substantially larger than ours, at 196-921 μmol m⁻² hr⁻¹. Likewise, Voynova et al., 2019 estimated net spring-summer sediment-water TA fluxes in the southern North Sea to be more than two orders of magnitude higher, at 488-1117 μmol m⁻² hr⁻¹ (Table 1 in Voynova et al., 2019). More in line with our findings are the results of Brenner et al., 2016, who directly measured TA fluxes in ex-situ incubations of 237.5 - 275 μmol m⁻² hr⁻¹ (Southern North Sea, 2011 and 2012), and a modelling simulation placing mean TA fluxes at 83 μmol m⁻² hr⁻¹ (Pätsch et al., 2018).

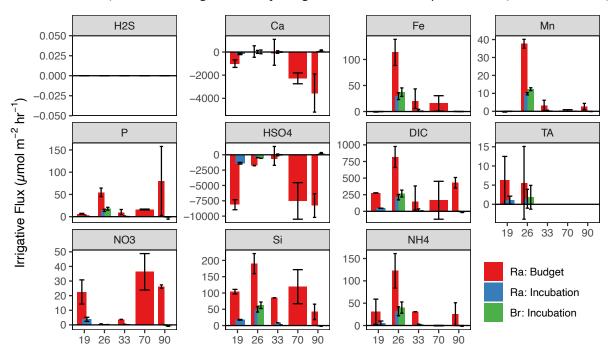


Figure 3. Average sediment-water fluxes (μmol m⁻² hr⁻¹) derived from Ra or Br incubation or Ra-decay from the North Sea sites only. In this figure, the error bars represent minimum and maximum fluxes calculated from replicate pore-water cores taken at each site.

3.2.2 Baltic Sea TA fluxes from PROFILE modelling

In the Baltic Sea muddy sites (AB, LB, MB), TA concentration in surface sediments (0-10 cm depth) exceeded surface water TA by 1-10 mM. This excess TA caused modelled sediment-water TA fluxes in the Baltic Sea muddy sites as high as 159 µmol m⁻² hr⁻¹ (site LB, Figure 4). Down-core trends in TA at these sites are well within measurement precision, allowing us to discuss potential sources of this TA and implications in section 3.5. In contrast with the relatively large excess TA in surface sediments at the Baltic Sea muddy sites, TA was very close to the overlying water concentration in Baltic sandy and



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silty sites, where increases in porewater TA were not observed until depths of 10-15 cm. Resulting modelled TA flux was small and negative at the Baltic sandy and silty sites, at -4.3 and -35 μ mol m⁻² hr⁻¹ respectively (Table 2 and Figure 4). In contrast, TA fluxes at the muddy sites were $53 \pm 95.3 \,\mu$ mol m⁻² hr⁻¹ (Table 2). This is approximately 50% of the non-resolved "nonriverine" TA source from Gustafsson et al., (2014), which was implied to be driven largely by denitrification, sulfur metabolism, and silicate weathering. The minor down-core differences in TA in the Baltic Sea sandy sites are due to enhanced mixing in sediments with relatively low microbial activity. An analysis (Sections 3.4-3.6), supporting the following factors as key in sustaining low or negative net TA fluxes in sandy Baltic Sea sites: 1) low net denitrification, 2) complete re-oxidation of sulfide and Fe(II) in surface sediments.

3.3 Sediment-water fluxes of other constituents (Baltic and North Sea)

Net sediment-water fluxes are shown together in Figure 4, with the bars representing average PROFILE-modelled fluxes, while the fluxes derived from incubation experiments and ²²⁴Ra decay balances are depicted as the colored points. Typically, PROFILE-modelled fluxes were at least 10x smaller than fluxes derived from the ²²⁴Ra decay balance. This is consistent with enhanced advective forcing that is captured by the decay-balance approach, but missed by the incubation experiment and PROFILE model, which are in closer agreement. In addition to the sediment-water fluxes described above, the PROFILE model also generates estimates of net internal production/consumption, representing a superimposition of combined biological and abiotic sources/sinks in each model zone of the sediment (Figure S2). By convention, positive irrigation or production values indicate a net increase in concentration.



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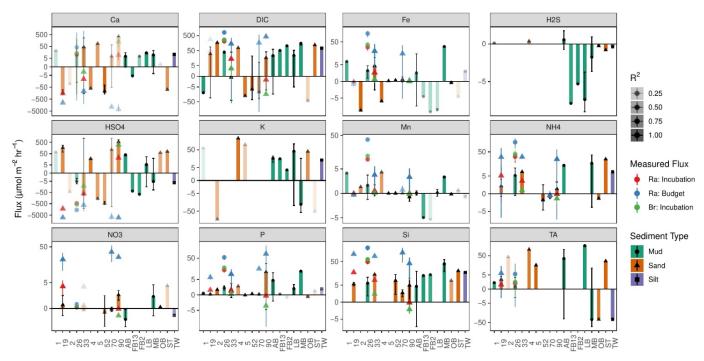


Figure 4. Sediment-water fluxes (μ mol m⁻² hr⁻¹) for all sites in Baltic (sites AB – TW) and North Seas (sites 1 – 90), derived from PROFILE modelling (colored bars), with incubation and decay rates for comparison from the North Sea only (colored points). Bars with greater transparency indicate PROFILE flux results of relatively lower confidence (R²). The error bars in this figure represent the standard deviation of fluxes modeled from replicate cores at each site.

Some solute fluxes exhibited similar trends across all sites in the North and Baltic Seas. For example, Si fluxes were consistently positive (out of sediments) for all sites, across all methods (Figure 4). In agreement with prior work in the region (Lipka et al, 2018; Gogina et al., 2018) ammonium (NH₄) release was also common and positive across all locations, except for three "sandy" sites in the North Sea (52, 70, 90). Net H₂S fluxes were not significantly different from zero in any North Sea sites, while PROFILE-modelled H₂S fluxes for three Baltic sites (FB2, FB13, and LB) were negative (into the sediment). We believe that these few large negative H₂S fluxes were created artificially by our use of a boundary condition (flux out of the bottom) that intersects with the zone of peak H₂S accumulation. In contrast, other modelled solute fluxes varied more between basins (North vs Baltic) than across sites within basins. For example, TA fluxes in the North Sea sites were close to zero (albeit with low R²), while PROFILE-modelled fluxes in the Baltic were larger in magnitude and more variable (Figure 4). Baltic sites were also more often sources of Ca to the water (except AB and FB13), while Ca fluxes in the North Sea were more variable and statistically less robust.

PROFILE-modelled NO₃ fluxes are variable and close to zero across all sites, in contrast to prior studies in the region suggesting net NO₃ uptake by denitrification (Brenner et al., 2016). In fact, net NO₃ fluxes derived from the incubation and Ra



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budget were positive (out of the sediment). This is in line with porewater NO₃ trends, which generally show no net depletion in porewaters relative to surface water. While canonical denitrification is likely playing some role, the small NO₃ fluxes in combination with larger NH₄ fluxes suggest that Dissimilatory Nitrate Reduction to Ammonium (DNRA) may be a larger sink for NO₃, and that any denitrification is fed by internally produced NO₃ rather than allochthonous NO₃. Therefore, in contrast with prior studies in the region (Burt et al., 2014; Brenner et al., 2016; Voyanova et al., 2019), we suggest a very limited potential for net TA production via denitrification in coastal North and Baltic Sea sediments. This is in line with De Borger et al (2021) and Pätsch et al (2018), who together indicate relatively low denitrification rates roughly balanced with nitrification, a feature that is supported by our low or negative modelled net TA fluxes (Figure S2).

DIC flux was variable across sites, but most often positive. Fluxes parameterized by Ra budgets and incubations agreed well with previously modelled diffusive fluxes for the region (Lipka 2017, figure 70), but our PROFILE model generated relatively low DIC fluxes. Sites with greater DIC release were also generally larger sources of Si, NH₄, and P, in line with the breakdown of algal detritus as a source for both.

3.4 Biogeochemical sources and sinks

3.4.1 Miller-Tans plots

The stable isotope composition of DIC reflects the diagenetic impact from dissolved organic matter (DOM) or methane oxidation and interactions with carbonates (e.g., Ku et al., 1999; Meister et al., 2019; Wu et al., 2017; Liu et al., 2021). Using measured δ^{13} C-DIC data, we conducted a Miller-Tans plot analysis aimed at identifying isotopic endmembers as possible DIC sources (Figure 5). In this graphical approach, the isotopic composition of DIC multiplied by its concentration is plotted against the concentration, with the slope indicative of the δ^{13} C-DIC source value. If the remineralization of particulate organic carbon (POC) is the only source of DIC to porewaters, the slope of the Miller-Tans plot should be the same as the isotopic signature of PO¹³C, which in these sites ranges from -20 to -25 % throughout the southern North Sea (Böttcher et al., 1998, 2000; Pollmann et al., 2021; Serna et al., 2014), from -21 to -24 % in the south-western Baltic Sea (Böttcher et al., unpublished data, Voß and Struck, 1997). Any deviation in the y-intercept from typical PO¹³C values may result from calcium carbonate dissolution as an important modulator of DIC and TA fluxes (Winde, 2017).

Muddy or silty sites (1, 26, AB, FB-13, FB-2, LB, MB, TW) tend to have a larger range in DIC and so allow for a more robust graphical analysis. δ^{13} C-DIC endmembers for these sites were generally much higher (heavier) in comparison with the PO¹³C references listed above, ranging from -16 to -21 ‰ (Baltic Sea) and -6 to -26 ‰ (North Sea). Due to lower microbial activity (Al Raei et al., 2009) and enhanced porewater exchange (de Beer et al., 2005), the sandy sites have a smaller total range in DIC concentrations and isotope compositions, challenging the application of a Miller-Tans Plot approach. Some of the sandy sites (2, 33, 4, OB, and ST), however, yield a similar δ^{13} C-DIC endmember value when compared to the muddy sites with values ranging between -16 and -19 ‰. Other sandy sites (19, 5, 52, 70, 90) have much heavier endmembers, and ranging from -6 to -12 ‰. This could suggest carbonate dissolution is contributing to heavier DIC values (Ku et al., 1999; Winde et



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al., 2014a,b; Wu et al., 2017). Carbonate weathering in pore waters has previously been suggested to be an important DIC and TA source (Winde et al., 2014a,b; Winde, 2017; Pätsch et al., 2018; Brenner et al., 2016). However, evidence for this is weak in our sandy North Sea sites, where porewater Ca concentrations were low relative to overlying surface water, as seen in the generally negative Ca fluxes (Figures 3 and 4), indicating no significant net carbonate dissolution. That the δ^{13} C-DIC endmember at Site 1 in the North Sea (-26 ‰) is well below the PO¹³C source indicates a potential DIC input from methane oxidation. Elevated surface water methane concentrations have been previously observed near Site 1 in the region around Helgoland (Bussmann et al., 2021).

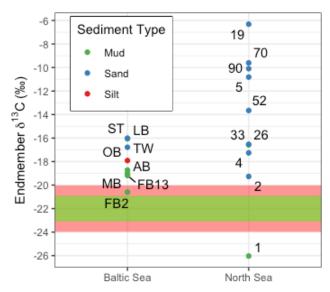


Figure 5. Potential end-member isotopic values for porewater DIC, derived from Miller-Tans plots of DIC * ¹³C-DIC vs DIC (shown in Fig S1). The colored areas represent a range in expected PO¹³C values for the North (red) and Baltic (green) seas.

3.4.2 DIC sources and sinks: Process correlations

While the ¹³C endmember indicated by the Miller-Tans plot approach above may help to identify possible electron donors, it cannot provide information as to the terminal electron acceptors that were used to respire the organic carbon substrate. Microbial sulfate reduction (MSR) is an important mineralization process in marine sediments of this region (Jörgensen, 1989; Al-Raei et al., 2009) and leads to the production of DIC and TA (Zeebe & Wolf-Gladrow, 2001). Downcore correlations between DIC and SO₄²⁻ may indicate the relative extent by which MSR is associated with DIC (and TA) production. We calculated excess DIC and sulfate as the difference between porewater and either 1) bottom water measurements, when porewater salinity, as inferred by porewater K concentration, did not change, or 2) from an empirical relationship between water-column DIC or SO₄²⁻ and K (Baltic sites FB-2 and FB-13, where salinity increased down-core).



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The muddy sites in the Baltic Sea have the strongest covariation of SO₄ deficit and DIC, with slopes ranging from -1 to -2.6, with r² between 0.86 – 0.97, suggesting MSR is a key anaerobic pathway of DIC formation in these sites (Figure 7, green points). However, Baltic Sea sandy sites and all North Sea sites (including sites 1 and 26, which are categorized as muds) have slopes closer to zero. In the sandy North Sea sites, SO₄²⁻ variation is larger than for DIC, suggesting calcification as a possible sink for DIC. In most cases, SO₄²⁻ concentrations are less than bottom water, indicating net consumption by MSR (19, 4, 5, 70, 90). However, there are also sites where some SO₄²⁻ measurements were greater than bottom water (52, 33), suggesting the oxidation of H₂S or FeS_x. This analysis is consistent with aerobic respiration as the major metabolic pathway for net DIC production in ventilated sandy sediments, likely supported by the relatively deep oxygen penetration depths.

Interestingly, North Sea mud stations (1 and 26) also have SO₄²:DIC slopes close to zero (-0.3 and 0.009, respectively). Therefore, MSR does not appear to be a dominant net DIC source in these muds, even though oxygen penetration depths are low and gross MSR rates are relatively high (Jørgensen, 1989). Instead, it is likely that MSR is coupled with internal sulfide oxidation. Potential sources of this sulfate in cases where it appears to be produced (North Sea sites 1, 33, and 52, and Baltic site ST) include sulfide oxidation and FeS_x oxidation, both of which would be increased by intermittent deeper oxygen penetration. The potential sink for DIC can be better understood when considering the TA:DIC ratios in the next section.

3.4.3 TA sources and sinks: Process correlations

We can directly assess the effect of metabolic processes on net TA production for a subset of sites where porewater TA was measured with a sufficient analytical precision. In most of the sandy sites, the down-core variation in TA was less than the analytical precision of the small-volume titrations used. For the sites where porewater TA data are available, TA and DIC are strongly correlated, even in some sandy sites with a much smaller range in both analytes (Figure 6, orange points). Muddy sites in the Baltic Sea (AB, LB, MB, and the silty TW) have a larger range in both TA and DIC, with slopes (i.e., ratio of TA:DIC) between 0.78 and 1.095. The two sandy Baltic sites (OB and ST) have similar slopes (1.16 and 0.76 respectively), although the range in both variables is much smaller. North Sea Site 1 (muddy) has a slope of 1.12, while the sandy sites have a broad range of 0.13 (site 2) to 0.65 (site 5) and 0.8 (site 4). Modelled DIC fluxes were much larger than TA fluxes for all North Sea sites (Figure 4; see scale), consistent with measured porewater TA:DIC slopes below 1:1 (Figure 6; orange points).

This is in part due to the likely dominance of aerobic respiration in ventilated North Sea sediments, which in contrast with anaerobic respiration, produces no TA. This dominance of aerobic respiration in permeable North Sea sediment can be attributed to the 'redox seal', which describes the situation in percolated sediment with mobile bed forms, where virtually the complete percolated sediment layer is oxygenized, whereas transport in the anoxic layer below is restricted to molecular diffusion (Ahmerkamp et a. 2015). Since porewater advection is several orders of magnitude more efficient in the transport of substrates and products than molecular diffusion, fluxes across the sediment surface sustained by aerobic process substantially exceed those sustained by anaerobic processes. Furthermore, we observe relatively large apparent SO₄²⁻ consumption below the oxic-anoxic interface (figure 6, green points), suggesting significant MSR even in sandy sediments. However, excess SO₄²⁻



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and DIC are not always positively correlated (figure 6), meaning that the DIC produced by MSR-driven DIC may be consumed insitu. One potential sink for this MSR-driven DIC production might be CaCO₃ precipitation in the oxic zone, which is densely colonized by calcifying infauna.

Trends between excess TA and apparent SO_4^{2-} consumption are like those for excess DIC (Figure 6; purple points). Baltic muddy sites have significant ($r^2 > 0.9$) slopes ranging from -1.3 to -2.1. Baltic sandy sites have slopes closer to zero, with small excess TA values, which appear disconnected from apparent SO_4^{2-} reduction. North Sea sandy sites also have slopes closer to zero, and the muddy site (Site 1) has a much shallower slope of -0.3.

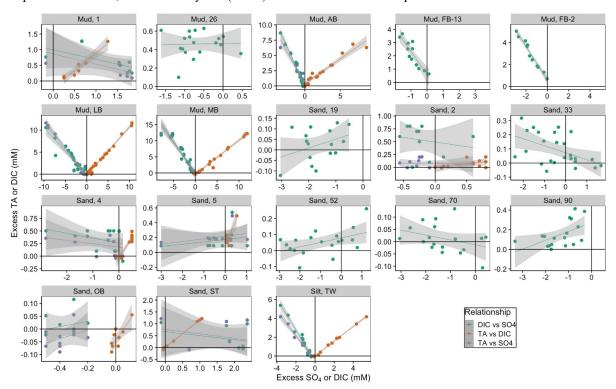


Figure 6. Relationships between excess DIC and excess SO₄²⁻ (green points), excess TA and excess DIC (orange), and excess TA vs excess SO₄²⁻ (purple). Positive values indicate a larger concentration in porewater relative to surface water.

3.5 PCA and regional patterns

To assess broader regional patterns in the drivers of sediment-water fluxes, we constructed a PCA (FactoMineR, Lê et. al. 2008) using a variety of selected parameters (Table 3). We included a set of biogeochemical parameters that appeared important in the above "process correlation" section, which included R_{DIC/SO4} (RDS) and the ratio of P v. Fe (RFeP), although excluded relationships with TA as that parameter was not available for the entire dataset. We also included the peak concentrations of Fe, sulfide, DIC, and Si (MaxFe, MaxHS, etc) and the depth at which this was observed (DFe, DHS, etc), as



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390 well as the first depth where sulfide was > 1 μM (DHS_1). The goal of this PCA was to identify general regional patterns in biogeochemical sources/sinks of TA and to relate them to readily available sediment characteristics.

Table 3. Input parameters used in the PCA analysis, and their definitions

Parameter	Definition
Phi	Porosity
RDS	Ratio change in DIC (mM) to change in Sulfate (mM)
RFeP	Ratio P (μ M) to Fe (μ M). Only slopes with $r^2 > 0.7$ are
	provided, others are marked as zero
MaxFe	Maximum porewater Fe (μM)
DFe	Depth at which Fe maximum occurs (cm)
MaxHS	Maximum porewater Sulfide (μM)
DHS_1	Depth where porewater Sulfide first exceeds 1 µM (up to
	total length of core)
MaxNO3	Maximum porewater NO3 or NOx concentration (μM)
DNO3	Depth at which maximum NO3 concentration occurs (cm)
MaxSi	Maximum porewater Si concentration (μM)
MaxDIC	Maximum porewater DIC concentration (mM)
MillerTans	slope of the Miller Tans plot

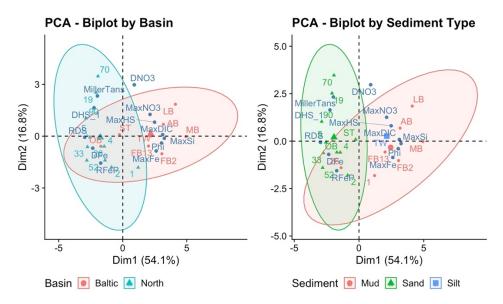


Figure 7: PCA Biplots separated by basin (left) and sediment type (right). Parameters (Table 3) are in dark blue, and the ellipses are calculated based on the sites, as separated by either basin (left) or sediment type (right).

Sites are strongly sorted by basin and sediment type along Dimension 1, which explains 54.8% of the variability (Figure 7). This is in part because the North Sea sites are dominated by sandy sites, and the Baltic sites by muddy sites. As a result, we also see strong sorting along the same Dimension 1, associated with the biogeochemical consequences of variable oxygen



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penetration driven by the interaction between porosity and advective porewater exchange. These factors include: 1) porosity, as low porosities can increase oxygen penetration with disturbance from currents, waves, and storms, 2) RDS and corresponding maximum HS, which indicates the importance of MSR, an anaerobic process, 3) Depth of max Fe, which accumulates as Fe(II) in anoxic porewaters and also is formed from the reduction of reactive Fe(III) minerals which (due to rapid (re)oxidation) accumulate at the oxic/anoxic boundaries. Additional sorting variables along Dimension 1 are related to organic matter accumulation, remineralization, and flushing of porewaters due to advection (North Sea) and include the dissolved nutrients and MaxFe. Dimension 2 explains 16.7% of the variability. Key sorting variables include the depth of Max NO₃, the Miller Tans slope, the accumulation of sulfide above 1 μ M, and the ratio of dissolved P to Fe. For example, North Sea sites with the highest Miller Tans slope (70, 19, 90 with potential δ^{13} C-DIC sources between -6 to -11‰) are also the North Sea sites with no significant relationship between dissolved P and Fe. Together, this supports the importance of MSR in mud/silt sediments, and O₂-based re-oxidation of reduced sulfur species following advective exchange in sandy sites with low organic matter inputs.

4. Conclusion

We combined surface water and porewater ²²⁴Ra measurements to build a decay balance, which indicates that the irrigative exchange between porewater and surface water is much greater than previously thought. Implicit in this assessment is our finding that the porewater ²²⁴Ra endmember was more variable and ~twice as large as previously assumed, of importance for future Ra-based budgets, which are highly sensitive to endmember quantification (Cook et al., 2018; Garcia-Orellana et al., 2021).

In our southern North Sea sites, net sediment-water TA fluxes were very small in comparison with previous estimates in this region (Voyanova et al., 2019; Burt et al., 2014). This is also in contrast to the measured and inferred TA production rates in the Wadden Sea, which are known to be 3-5 orders of magnitude higher on an aerial basis (Santos et al., 2015), largely associated with substantial net MSR during warm seasons (Al-Raei et al., 2009). TA in ventilated North Sea surface sediments was similar to surface water concentrations, causing net sediment-water fluxes to also be very low (~10x less than for DIC). Here, TA sources and sinks are closely balanced in North Sea sediments. We ascribe this to the potential influence of 1) DNRA / denitrification being fed internally by recycled, rather than "new" NO₃, 2) MSR in close balance with sulfide oxidation, and 3) carbonate mineral precipitation in the oxic zone. While positive Si fluxes could suggest silicate weathering as TA source, we cannot yet ascribe this to the erosion of biogenic (e.g., diatoms) or geogenic minerals.

In the Baltic Sea sites, excess TA was greater in muddy regions (and close to equilibrium in silty and sandy sites), causing modelled fluxes to be generally low and positive (muddy) or negative (sands) implying a sink in surficial sediments. At Baltic Sea sites with small positive TA fluxes (muds/silt), net TA production is likely due to net MSR (see sulfate consumption; Figure 7) and associated pyrite burial. In contrast, at sites with net modelled TA uptake (sands), modelling results indicate TA consumption in the uppermost sediment layers, which we attribute to: 1) re-oxidation of sulfide and Fe(II), 2) minimal net denitrification, as in North Sea sites, and 3) carbonate mineral dissolution.





Our findings suggest that coastal sediments of the Southern North Sea and Western Baltic Sea are currently not major sources of TA to the water column. This is because TA production in deeper anaerobic sediments is counter-balanced by reoxidation and TA consumption in overlying oxic sediment, causing net TA fluxes to be small and variable. The seasonality of these TA fluxes and their impact on water-column carbonate chemistry and ultimately air-water CO₂ exchange was not assessed in this study but should be a topic of future research.

440 Data Availability

Upon publication, all data used in the preparation of this manuscript will be made publicly available at the Helmholtz Coastal Data Center (HCDC), an open-access repository, according to the FAIR principles.

Author Contribution

Contributor roles according to the CRediT Taxonomy

445 Conceptualization: BVD, HT, MEB

Formal Analysis: BVD, MAZ, AN

Investigation, inclusive of field and laboratory work: BVD, NL, MAZ, AN, ML, DP

Project Administration, Resources, Supervision: HT, MEB, DP

Writing - original draft preparation: BVD, MAZ

450 Writing – review and editing: BVD, NL, MAZ, AN, DP, HT, MEB

Competing interests

The authors declare that they have no conflict of interest.

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