

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

5 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

⁴ Alfred 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

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

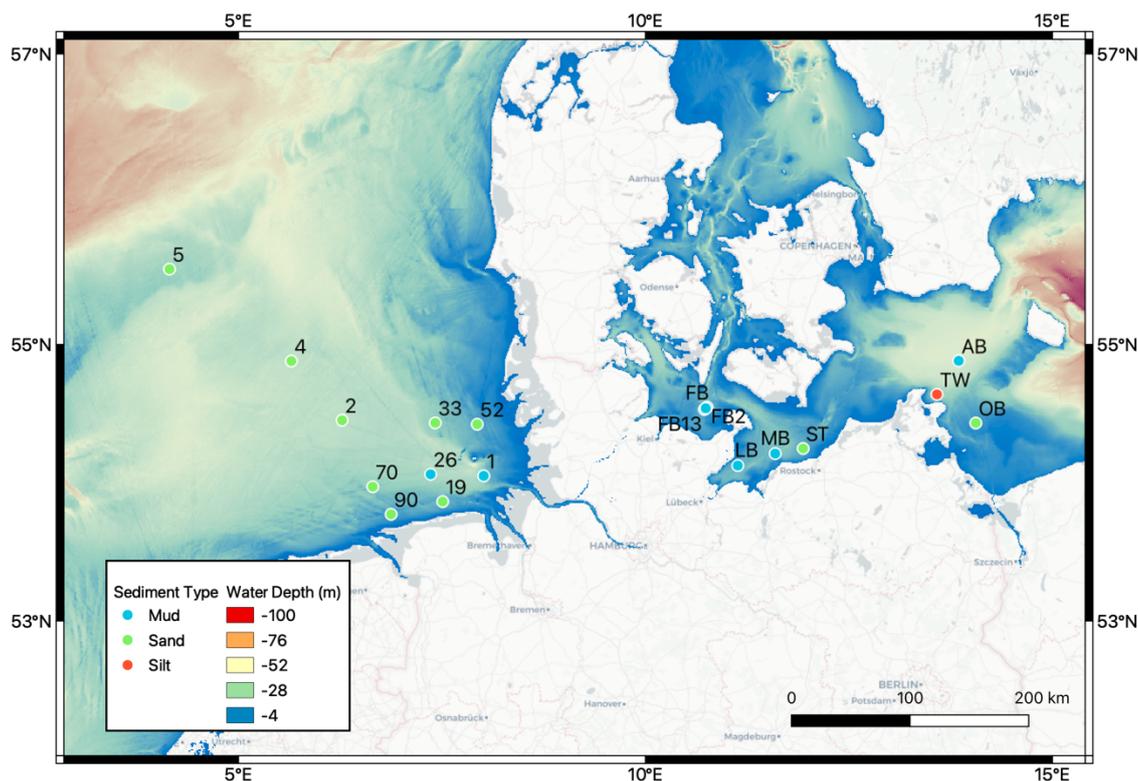
15 **Abstract.** Benthic alkalinity production is often suggested as a major driver of net carbon sequestration
in continental shelf ecosystems. However, information and direct measurements of benthic alkalinity
fluxes are limited and are especially challenging when biological and dynamic physical forcing 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) ²²⁴Ra budgeting, 2)
20 incubations with ²²⁴Ra 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
25 previously estimated for the region, in part due to measured porewater ²²⁴Ra activities higher than
previously assumed. Net alkalinity fluxes in the Baltic Sea were relatively low, ranging from an uptake
of -35 $\mu\text{mol m}^{-2} \text{hr}^{-1}$ to a release of 53 $\mu\text{mol m}^{-2} \text{hr}^{-1}$, and in the North Sea from 1 to 33.6 $\mu\text{mol m}^{-2} \text{hr}^{-1}$.
Lower than expected apparent nitrate consumption (potential denitrification), across all sites, is one
30 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

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
35 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 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
40 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
45 (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.,
50 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.
55 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
60 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.



65 **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 finer-grained 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
90 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
95 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
100 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 SO_4^{2-} , DIC, TA, Fe, and PO_4^{3-} 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.,
105 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 CO_2 uptake and, therefore, require further
110 investigations.

2 Methods

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
115 (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
120 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 \pm S.D.), with a maximum 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).

125 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 μm) 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.

130 2.2 Lab Analyses

The samples taken during the HE541 cruise were analyzed according to the following protocols: TA
was determined at the Leibniz-Institut für Ostseeforschung Warnemünde (IOW) via a small volume
(400-800 μL), two-point potentiometric titration according to van den Berg and Rogers, 1987. The
sample was transferred into a vial with a pre-set volume of HCl on board, assuming that all weak acids
135 in solution would be protonated ($\text{pH} < 3$). The sample was cooled at 4 $^{\circ}\text{C}$ 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 μL , 0.1 M), the second point was recorded ($\text{pH} \approx 2$). Taking these two points, the slope of the
electrode response was determined (van den Berg and Rogers, 1987). The measurements were
140 calibrated against certified reference materials (CRMs), from batch 142 provided by Prof Andrew
Dickson (Scripps Institution of Oceanography, San Diego, USA).

DIC and $\delta^{13}\text{C}$ -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,
145 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).

150 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,
155 United Kingdom) with the addition of Fe and Mn ($c = 1 \text{ mg/L}$). 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 as described by Winde et al. (2014a) and Lipka
160 et al. (2018).

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
165 fluxes span a complete range in possible rates.

First, a water-column ^{224}Ra 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 ^{224}Ra employed as tracers. We also used PROFILE
170 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).

2.3.1 ^{224}Ra decay balance

Samples for ^{224}Ra 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 (^{224}Ra determinations were not made
175 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
180 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 ^{224}Ra 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.

185 Because ^{224}Ra 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 ^{228}Th), 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
190 between radioactive decay and the measured surface water activity. These irrigation rates derived from
the ^{224}Ra decay balance (mean = $81 \text{ L m}^{-2} \text{ d}^{-1}$) 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
195 was used for the "muddy" and "silty" sites. Sign convention is that positive fluxes indicate a flux out of
the sediment and vice versa.

2.3.2 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
200 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 nutrient and Bromide analyses 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 µm syringe filters and stored frozen until analysis in the land-based laboratory. After incubations, additional samples from the supernatant were immediately taken for measurements of Radium and DIC. The incubation method is described in detail in Neumann et al. (2021).

2.3.3 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. Biodiffusivity and irrigation coefficients were chosen through an iterative process, changing one variable at a time and comparing model performance for each sediment class.

Table 1. Parameterization of PROFILE model

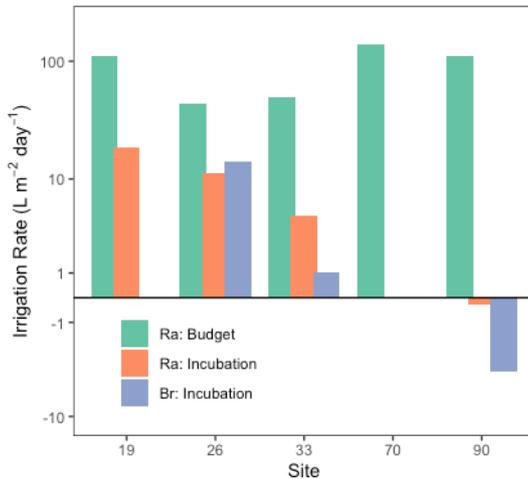
		Sand	Silt	Mud
Max deviation when accepting value (%)		0.001		
Level of significance for F-statistic		0.2		
Biodiffusivity (cm² hr⁻¹)	< 5 cm	0.36	0.18	0.18
	5-10 cm	0.18	0	0
	>10 cm	0	0	0
Irrigation coefficient (hr⁻¹)	< 5 cm	0.11	0.0011	0.0011
	5-10 cm	0.011	0	0
	>10 cm	0	0	0

Porosity (φ)	0.4	0.6	0.8
Coefficient of Molecular Diffusivity ($\text{cm}^2 \text{hr}^{-1}$)	Calculated in R: 'marelac' as $f(\text{Sal}, \text{Temp}, \text{Pres})$		
Sediment Diffusivity ($\text{cm}^2 \text{hr}^{-1}$)	$D_s = D / (1 + 3[1 - \varphi])$		
Boundary Conditions	Concentration and diffusive flux at the bottom		

3 Results and Discussion

3.1 North Sea Irrigation rates (decay balance and incubation)

235 Surface water ^{224}Ra activities were consistent throughout the southern North Sea, ranging from
248 $\pm 11.6 \text{ dpm m}^{-3}$ (Site 33) to $384 \pm 21.3 \text{ dpm m}^{-3}$ (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
240 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 $44 \pm 3.0 \text{ dpm L}^{-1}$; average of 19.9 dpm L^{-1} (or $19,946 \text{ dpm m}^{-3}$)) 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 = $81 \text{ L m}^{-2} \text{ d}^{-1}$
245 $^{-1}$; 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 44 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).



250 **Figure 2.** Porewater irrigation rates ($\text{L m}^{-2} \text{d}^{-1}$) for the North Sea only, derived from Ra budget (green) and incubations with Ra (orange) or Bromide (blue) as tracers.

255 Irrigation rates derived from Br and ^{224}Ra measurements during the ship-board incubation
 experiments (mean = $10.5 \text{ L m}^{-2} \text{ d}^{-1}$; blue bars) were a factor of 8 lower than rates derived from the
 ^{224}Ra 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
 260 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
 265 varying importance of physical and bio-mediated forcing of net sediment-water exchange.

3.2 Net TA Fluxes

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,
 270 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.,
 275 2019). Excluding the PROFILE-modelled TA fluxes for the North Sea (which had very low R^2), net TA
 fluxes in the North Sea were relatively low, ranging from 1 to $6.3 \mu\text{mol m}^{-2} \text{ hr}^{-1}$ across all sites and
 sediment types (Table 2).

280 **Table 2.** Comparison of average TA fluxes ($\mu\text{mol m}^{-2} \text{ hr}^{-1}$) 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 ± 43.5 (n=2)	-35 (n=1)
North Sea	PROFILE	1 ± 2.1 (n=3)	33.6 ± 42.3 (n=5)	-
North Sea	Ra Incubation	1.5 ± 3.5 (n=2)	1.1 ± 1.5 (n=2)	-
North Sea	Ra Budget	5.6 ± 13.4 (n=2)	6.3 ± 8.8 (n=2)	-
North Sea	Br Incubation	1.8 ± 4.4 (n=2)	-	-

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
 285 (0.002 – 0.004 mM) greater than bottom water TA, an insignificant difference considering the relatively
 low analytical precision of these small-volume TA titrations. As a result, the inferred sediment-water

TA fluxes in Burt et al., 2014 were substantially larger than ours, at $196\text{--}921 \mu\text{mol m}^{-2} \text{hr}^{-1}$. Likewise, Voynova et al., 2019 estimated sediment-water TA fluxes in the southern North Sea (based on observed seasonal variations in TA) to be more than two orders of magnitude higher, at $488\text{--}1117 \mu\text{mol m}^{-2} \text{hr}^{-1}$ (Table 1 in Voynova et al., 2019). Two recent modelling and empirical studies placed mean TA fluxes for this region at $83 \mu\text{mol m}^{-2} \text{hr}^{-1}$ (Pätsch et al., 2018) and $238\text{--}275 \mu\text{mol m}^{-2} \text{hr}^{-1}$ (Brenner et al., 2016) respectively, closer to but still greater than our results.

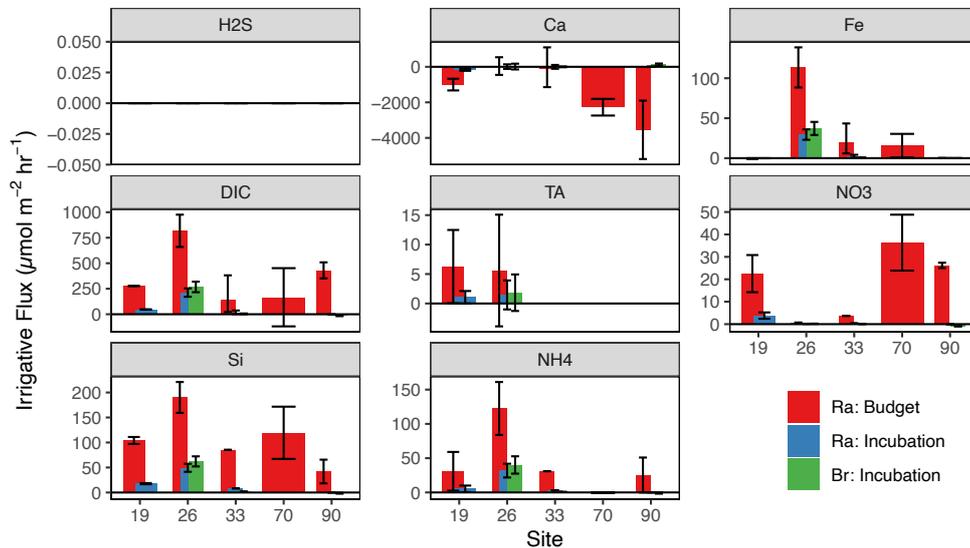


Figure 3. Average sediment-water fluxes ($\mu\text{mol m}^{-2} \text{hr}^{-1}$) derived from Ra or Br incubation or Ra-decay from the North Sea sites only. In this figure, the error bars represent minimum and maximum fluxes calculated from replicate pore-water cores taken at each site.

3.2.2 Baltic Sea TA fluxes from PROFILE modelling

In the Baltic Sea muddy sites AB and LB, TA concentration in surface sediments (0-10 cm depth) exceeded surface water TA by 1-10 mM, a significant excess considering the analytical precision of small-volume TA titrations. This excess TA caused PROFILE-modelled sediment-water TA fluxes in the Baltic Sea muddy sites AB and LB as high as $159 \mu\text{mol m}^{-2} \text{hr}^{-1}$ (site LB, Figure 4), while a smaller negative flux was observed at site MB. 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 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 $-35 \mu\text{mol m}^{-2} \text{hr}^{-1}$ respectively (Table 2 and Figure 4). In contrast, PROFILE-modelled TA fluxes averaged across all Baltic Sea muddy sites for which fluxes could be calculated (AB, LB, MB) were $53 \pm 95.3 \mu\text{mol m}^{-2} \text{hr}^{-1}$ (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
 315 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 ^{224}Ra decay balances are depicted as the colored points. Typically, PROFILE-modelled fluxes were at least 10x
 320 smaller than fluxes derived from the ^{224}Ra 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
 325 the sediment (Figure S2).

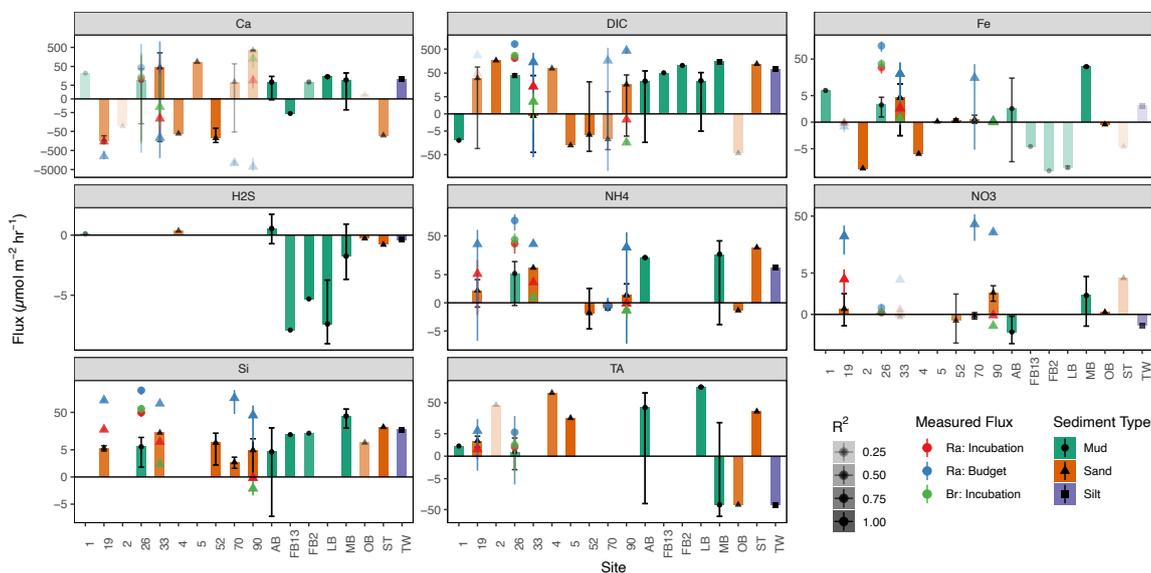


Figure 4. Sediment-water fluxes ($\mu\text{mol m}^{-2} \text{hr}^{-1}$) for all sites in Baltic (sites AB – TW) and North Seas (sites 1 – 90), derived from PROFILE modelling (colored bars), with incubation and decay rates for comparison from the North Sea only (colored points). Bars with greater transparency indicate PROFILE flux results of relatively lower confidence (R^2). The error bars in this figure represent the standard deviation of fluxes modeled from replicate cores at each site.
 330

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 (NH_4^+) release was also common and positive across all locations, except for three “sandy” sites in the North Sea (52, 70, 90). Net H_2S fluxes were not significantly different from zero in any North Sea sites,
 335

340 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).
345 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 budget were positive (out of the sediment). This is in
350 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 the presence of Dissimilatory Nitrate Reduction to Ammonium (DNRA) as a secondary sink for NO₃, with denitrification fed by internally produced NO₃ rather than allochthonous NO₃. Low net denitrification is consistent with slurry and flow-through core incubations
355 in the German Bight showing denitrification largely supplied internally by nitrification (Marchant et al., 2016). In contrast, net denitrification can be quite high at estuarine sites receiving “new” NO₃ from the Elbe river, especially during the winter when river discharge is elevated (Neumann et al., 2017). Nevertheless, DNRA is a relatively minor process in the German Bight, limited mostly to intermediate (Marchant et al., 2016) and fine-grained sediment (Bratek et al., 2020). As such, it could also be that
360 these NH₄⁺ fluxes are a simple result of aerobic degradation of organic matter. Therefore, in contrast with prior studies in the region (Burt et al., 2014; Brenner et al., 2016; Voyanova et al., 2019), we suggest relatively low net denitrification rates, with limited potential for TA production. 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
365 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
370 source for both.

3.4 Biogeochemical sources and sinks

3.4.1 δ¹³C-DIC sources: 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 δ¹³C-DIC data, we conducted a Miller-Tans plot analysis aimed at identifying isotopic endmembers as possible DIC sources (Figure 5). In this
375 graphical approach, the isotopic composition of DIC multiplied by its concentration is plotted against the concentration, with the slope indicative of the δ¹³C-DIC source value, or endmember. If the

380 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 PO^{13}C , which in these sites
ranging 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 PO^{13}C values
385 (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. Calculated $\delta^{13}\text{C}$ -DIC endmembers for these sites
were generally much higher (heavier) in comparison with the PO^{13}C references listed above, ranging
from -16 to -21 ‰ (Baltic Sea) and -17 to -26 ‰ (North Sea). That the $\delta^{13}\text{C}$ -DIC endmember at Site 1
390 in the North Sea (-26 ‰) is well below the PO^{13}C 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). 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, decreasing the confidence in a Miller-Tans Plot
395 approach. Some of the sandy sites (2, 33, 4, OB, and ST), however, yield a similar $\delta^{13}\text{C}$ -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 al., 2014a,b; Wu et al., 2017). Carbonate weathering in pore waters has previously been suggested to
400 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. Instead, these heavier $\delta^{13}\text{C}$ -DIC
endmembers are likely reflective of iterative precipitation/dissolution cycles (Walter et. al. 2007).

405

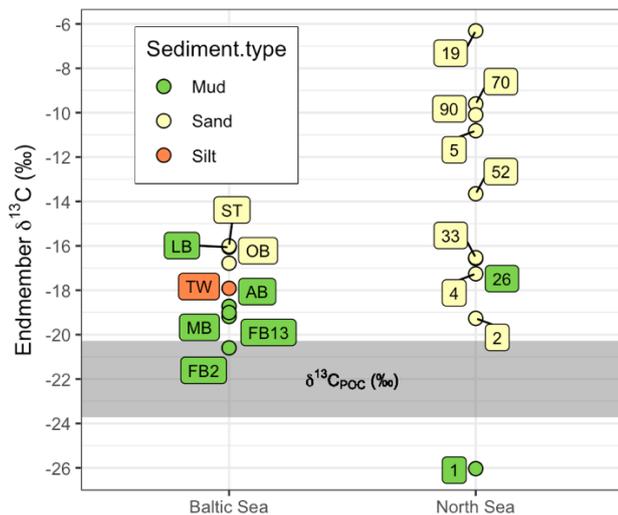


Figure 5. Potential end-member isotopic values for porewater DIC, derived from Miller-Tans plots of DIC * $\delta^{13}\text{C}$ -DIC vs DIC (shown in Fig S1). The grey area represents a range in expected PO^{13}C values for the North Baltic seas combined.

410 3.4.2 DIC sources and sinks: Process correlations

While the $\delta^{13}\text{C}$ -DIC source 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 (Jørgensen, 1989; Al-Raei et al., 2009) and leads to the production of DIC and TA (Zeebe & Wolf-Gladrow, 2001). Down-core correlations between DIC and SO_4^{2-} 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 SO_4^{2-} and K (Baltic sites FB-2 and FB-13, where salinity increased down-core).

The muddy and silty sites in the Baltic Sea (MB, AB, LB, TW) have the strongest covariation of SO_4 deficit and DIC, with slopes ranging from -1 to -2.6, with r^2 between 0.86 – 0.97, suggesting MSR is a key anaerobic pathway of DIC formation in sites with fine-grained sediment (Figure 7, green points). In sandy North Sea sites, SO_4^{2-} variation is larger than for DIC, suggesting calcification as a possible sink for DIC. In most cases, SO_4^{2-} concentrations are less than bottom water, indicating net consumption by MSR (19, 4, 5, 70, 90). However, there are also sites where some SO_4^{2-} measurements were greater than bottom water (52, 33), suggesting the oxidation of H_2S 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 DIC: SO_4^{2-} slopes close to zero (-0.3 and 0.009, respectively). At these sites, while SO_4 variation is greater than DIC variation, the net change in SO_4 is still low. This suggests that, even though oxygen penetration depths are low and gross MSR rates are relatively high (Jørgensen, 1989), 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.

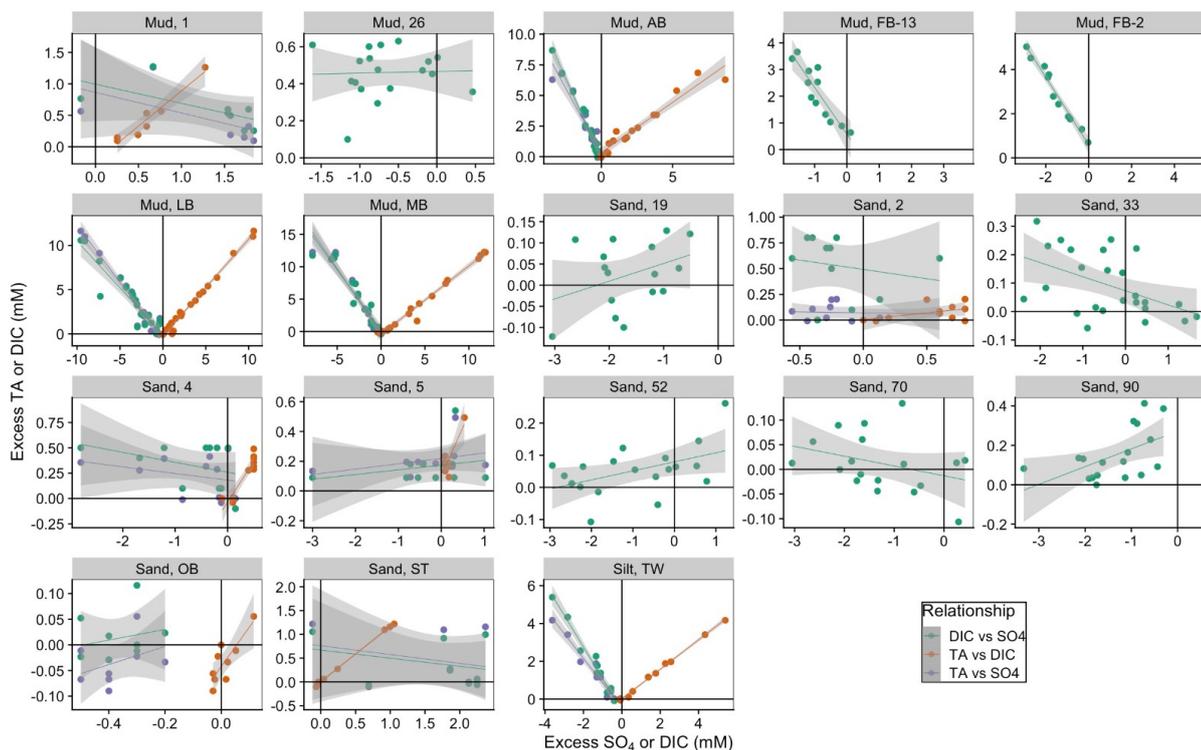
3.4.3 TA sources and sinks: Process correlations

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).

450 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
455 diffusion (Ahmerkamp et al. 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 SO_4^{2-} consumption below the oxic-anoxic interface
(figure 6, purple points), suggesting significant MSR even in sandy sediments.

460 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. We infer that iterative carbonate mineral dissolution
and precipitation recycling is responsible for our observations of relatively high TA:DIC ratios as well
465 as heavy $\delta^{13}\text{C}$ -DIC endmembers (Miller-Tans slope). Likewise, we infer microbial SO_4 reduction is also
matched by sulfide oxidation, with little net impact on sediment-water TA fluxes. Together, it appears
that strong internal recycling of carbonate minerals and SO_4 is a major factor causing net TA fluxes to
be low across all sites (Table 1).



470 **Figure 6.** Relationships between excess DIC and excess SO_4^{2-} (green points), excess TA and excess
 475 DIC (orange), and excess TA vs excess SO_4^{2-} (purple). Positive values indicate a larger concentration in
 porewater relative to surface water.

4. Conclusion

We combined surface water and porewater ^{224}Ra measurements to build a decay balance, which
 475 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 ^{224}Ra endmember was
 more variable and \sim 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).

480 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
 485 surface water concentrations, causing net sediment-water fluxes to also be very low (\sim 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”
 NO_3 , 2) MSR in close balance with sulfide oxidation, and 3) carbonate mineral precipitation in the oxic

490 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). In contrast, at sites with
495 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
500 anaerobic sediments is counter-balanced by re-oxidation 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.

Data Availability

505 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](#)

510 *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

515 *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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