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## Dark carbon fixation in the Arabian Sea oxygen minimum zone contributes to sedimentary organic carbon (SOM)

Sabine K. Lengger<sup>1,2,3\*</sup>, Darci Rush<sup>3</sup>, Jan Peter Mayser<sup>2</sup>, Jerome Blewett<sup>2</sup>, Rachel
 Schwartz-Narbonne<sup>4</sup>, Helen M. Talbot<sup>4,†</sup>, Jack J. Middelburg<sup>5</sup>, Mike S.M. Jetten<sup>6</sup>, Stefan
 Schouten<sup>3,5</sup>, Jaap S. Sinninghe Damsté<sup>3,5</sup> and Richard D. Pancost<sup>2</sup>

- <sup>6</sup> <sup>1</sup> Biogeochemistry Research Centre, School of Geography, Earth and Environmental Science,
- 7 University of Plymouth, PL48AA, Plymouth, United Kingdom.
- <sup>2</sup> Organic Geochemistry Unit, School of Chemistry, University of Bristol, BS81TS, Bristol,
   9 United Kingdom.
- <sup>3</sup> NIOZ Royal Netherlands Institute for Sea Research, Dept. of Marine Microbiology and
- Biogeochemistry, and Utrecht University, 1797SZ, Texel, The Netherlands.
- <sup>4</sup> School of Natural and Environmental Sciences, Newcastle University, Drummond Building,
- 13 NE1 7RU, Newcastle-upon-Tyne, United Kingdom.
- <sup>5</sup> Department of Earth Sciences, Faculty of Geosciences, Utrecht University, 3508 TA, Utrecht,
   The Netherlands.
- <sup>6</sup> Department of Microbiology, IWWR, Radboud University Nijmegen, 6525 XZ, Nijmegen, The
   Netherlands
- 18
- 19 \* corresponding author: Sabine Lengger (sabine.lengger@plymouth.ac.uk)
- <sup>20</sup> † present address: BioArCh, Environment Building, University of York, YO10 5DD, Heslington,
- 21 United Kingdom.

#### 22 Key Points:

- One fifth of organic matter on Arabian Sea seafloor could stem from bacterial carbon
   fixation in the oxygen minimum zone
- Evaluation of past anoxic events needs to take chemoautotrophic contribution into
   account in isotope balances
- Biogeochemical models ignoring dark carbon fixation could highly underestimate oxygen demand and thus expansion of oxygen minimum zones
- 29

#### 30 Abstract

In response to rising CO<sub>2</sub> concentrations and increasing global sea surface temperatures, oxygen 31 minimum zones (OMZ), or "dead zones", are expected to expand. OMZs are fueled by high 32 primary productivity, resulting in enhanced biological oxygen demand at depth, subsequent 33 oxygen depletion, and attenuation of remineralization. This results in the deposition of organic 34 carbon-rich sediments. Carbon drawdown is estimated by biogeochemical models; however, a 35 major process is ignored: carbon fixation in the mid- and lower water column. Here, we show 36 that chemoautotrophic carbon fixation is important in the Arabian Sea OMZ; and manifests in a 37 <sup>13</sup>C-depleted signature of sedimentary organic carbon. We determined the  $\delta^{13}$ C values of SOM 38 deposited in close spatial proximity but over a steep bottom-water oxygen gradient, and the  $\delta^{13}$ C 39 composition of biomarkers of chemoautotrophic bacteria capable of anaerobic ammonia 40 oxidation (anammox). Isotope mixing models show that detritus from anammox bacteria or other 41 42 chemoautotrophs likely forms a substantial part of the organic matter deposited within the Arabian Sea OMZ (~17%), implying that the contribution of chemoautotrophs to settling organic 43 matter is exported to the sediment. This has implications for the evaluation of past, and future, 44 OMZs: biogeochemical models that operate on the assumption that all sinking organic matter is 45 photosynthetically derived, without new addition of carbon, could significantly underestimate the 46 extent of remineralization. Oxygen demand in oxygen minimum zones could thus be higher than 47

48 projections suggest, leading to a more intense expansion of OMZs than expected.

#### 49 Plain Language Summary

50 Oxygen minimum zones are areas in the ocean in which algae produce large amounts of organic

51 material. When this sinks towards the seafloor, all oxygen at depth is used up. This results in vast

<sup>52</sup> "dead zones" where almost no oxygen is available to sustain life. With global warming, and

- increased nutrients from rivers, dead zones are forecast to expand. Computer models can
- calculate this, by considering algal production, and the amount of material delivered to the
- seafloor. However, these models often ignore a major process: anaerobic bacteria in the deeper

water column, that can live at the edge or in the middle of these dead zones, which can also

57 produce organic material from the dissolved  $CO_2$ . In this study, we used the fact that these

bacteria add a distinct signature to the organic material, to show that one fifth of the organic

59 matter on the seafloor could stem from bacteria living in these dead zones. Thus, models that

have missed out on considering this contribution could have underestimated the extent of oxygen

61 depletion we are to expect in a future, warming world. A more intense expansion of dead zones

than expected could have severe ecological, economical (fisheries), and climatic consequences.

63

#### 64 1 Introduction

Marine primary production fixes 50 Pg carbon per year, of which only about 1% is buried 65 in sediments (Dunne et al., 2007; Middelburg, 2011). The majority of organic carbon derived 66 from the photic zone is remineralised during sedimentation, fueling heterotrophic bacterial 67 activity in the water column (Keil et al., 2016). In marginal settings and OMZs, marine primary 68 production in the photic zone can be significantly higher than in other settings. Organic carbon 69 (OC) sedimentary accumulation rates within an OMZ can be in the range of tens to hundreds of 70 mg C cm<sup>-2</sup> y<sup>-1</sup> (Hartnett et al., 1998; Hedges and Keil, 1995) higher than observed in other parts 71 of the ocean. These high accumulation rates are most commonly attributed to attenuation in 72 remineralization rates within the OMZ, and low bottom-water oxygenation, which results in 73 decreased biodegradability of polymeric and matrix-protected substances (Burdige, 2007). 74

As a consequence of increasing atmospheric CO<sub>2</sub> concentrations and, consequently, temperature, oceanic OMZs are forecast to expand in a fashion similar to the past (Breitburg et al., 2018; Queste et al., 2018; Schmidtko et al., 2017; Shaffer et al., 2009; Stramma et al., 2010). Whilst the expansion of OMZs will result in widespread habitat loss of marine life and could cause an increase in emissions of greenhouse gases such as N<sub>2</sub>O and CH<sub>4</sub>, it could also act as a long-term negative feedback on global warming via the enhanced drawdown and storage of organic carbon in sediments.

The biogeochemical system in subsurface waters, where light does not penetrate, has 82 recently emerged to be substantially more complex – and possibly more important for the global 83 84 carbon cycle - than previously assumed. In particular, dark water-column microbial activity is higher than what can be accounted for by heterotrophs (Herndl and Reinthaler, 2013), suggesting 85 an important role for chemoautotrophy, i.e. fixation of dissolved inorganic carbon (DIC). It has 86 been suggested to contribute substantially to the global carbon budget, with estimates ranging 87 from 0.11 to 1.1 Pg C y<sup>-1</sup>, equating to ca. 2% of total estimated yearly marine primary production 88 (Middelburg, 2011; Reinthaler et al., 2010). The predominant chemoautotrophic process in the 89 oxic, dark, pelagic ocean is thought to be nitrification (Middelburg, 2011; Pachiadaki et al., 90 2017). When oxygen is limited, nitrification still occurs, but other chemoautotrophic processes 91 92 dominate, such as anaerobic oxidation of ammonia and sulfide oxidation (Ulloa et al., 2012; Wright et al., 2012). 93

Under hypoxic conditions, such as in the water column of OMZs, both archaeal (aerobic)
and anaerobic oxidation of ammonia are thought to dominate dark inorganic carbon fixation
processes (Lam and Kuypers, 2010; Pitcher et al., 2011). Here, nitrite accumulates, and other
anaerobic autotrophic processes such as sulfide oxidation and methanogenesis are suppressed,
most likely due to the abundance of nitrate and ammonia (Canfield, 2006; Ulloa et al., 2012).

99 Of the inorganic carbon converted to organic matter within the OMZ, only a negligible fraction is presumably transported to the sediments and preserved, as this newly produced 100 material is regarded as more labile than the sinking OC derived from the photic zone (Cowie and 101 Hedges, 1992; Keil et al., 1994; Middelburg, 1989). Dark carbon fixation rates are challenging to 102 quantify: they have been determined experimentally (Reinthaler et al., 2010; Taylor et al., 2001), 103 or have been estimated from the reaction stoichiometry of respiration based on Redfield organic 104 105 matter and growth yields of nitrifiers (Middelburg, 2011; Wuchter et al., 2006). In OMZs, such as the Peruvian margin (Lam et al., 2009), the Arabian Sea (Jensen et al., 2011), or the sulfidic 106 Black Sea (Lam et al., 2007), the activity of some chemoautotrophs was determined via <sup>15</sup>N-107

labelling, and formation of the products of their biogeochemical reactions. However, incubation
 methods may suffer from bias, because *in situ* conditions such as pressure are difficult to

110 maintain.

However, carbon from within anoxic waters has been observed in some settings to

112 contribute to the particulate OC flux: for example, in eutrophic lakes (Hollander and Smith,

- 113 2001) and anoxic fjords (van Breugel et al., 2005b). Furthermore, discrepancies between
- 114 modelled and observed organic carbon fluxes suggest that dark carbon fixation in anoxic marine
- settings significantly contributes to sinking material (Keil et al., 2016; Taylor et al., 2001).

One way to constrain this input into sedimentary organic matter is to use isotope mixing 116 models. Photosynthetically fixed carbon generally has stable carbon isotopic compositions of ca 117 -19 to -21 ‰ due to Rubisco fixation. However, chemoautotrophs dwelling in OMZs typically 118 have a lower  $\delta^{13}$ C values. This is the result of multiple factors: they either use  ${}^{13}$ C-depleted CO<sub>2</sub> 119 generated by remineralization, have larger fractionation factors due to the higher abundance of 120 CO<sub>2</sub> at depth (Freeman et al., 1994), or use carbon fixation pathways such as the acetyl 121 coenzyme A pathway, which results in <sup>13</sup>C-depleted biomass (Hayes, 2001). This characteristic 122 chemoautotrophic isotopic signature in the organic carbon could allow us to quantify the 123 contribution of dark carbon fixation to sedimentary organic matter. 124

Here, we investigated the  $\delta^{13}$ C value of sedimentary organic matter of surface sediments 125 deposited within the OMZ of the Arabian Sea and employed a simple isotope mixing model to 126 investigate the extent of input from OMZ carbon fixation into sedimentary organic matter. As the 127 128 major process in the Arabian Sea OMZ known to produce isotopically light biomass is anaerobic oxidation of ammonia (anammox; Ulloa et al., 2012; Villanueva et al., 2014), in order to 129 determine the isotopic signature of this pathway, we developed and applied a method to 130 determine the  $\delta^{13}$ C values of a novel biomarker, bacteriohopanetetrol stereoisomer (BHT'), 131 which has been found to be unique to anammox bacteria in the marine environment in culture 132 and environmental studies (Rush et al., 2014, 2019; Rush and Sinninghe Damsté, 2017). It has 133 been found in the Arabian Sea and other marine anoxic settings (Matys et al., 2017; Sáenz et al., 134 2011). We also used stable isotope probing experiments to exclude sedimentary anammox as an 135 136 important contributor to this process. This allowed us to investigate the contribution of these dark carbon fixers to sedimentary organic carbon. 137

#### 1382Materials and Methods

139 2.1 Sediment sampling and stable isotope probing incubations

Sediments were collected with multicore devices on the R/V Pelagia in the Northern 140 Arabian Sea in January 2009 during the PASOM cruise 64PE301 along the Murray Ridge (Fig. 141 1f), which protrudes into the core of the OMZ. Two cores, one each from P900 (885 m water 142 depth) and P1800 (1786 m water depth), hereinafter referred to as anoxic and oxic, respectively, 143 were incubated on board as described by Pozzato et al. (2013 a, b). In brief, particulate or 144 dissolved organic matter from the diatom *Thalassiosira pseudonana* containing 20 and 18 % <sup>13</sup>C, 145 respectively, were added to the tops of core tubes of 10 cm internal diameter. Between 2 and 6 % 146 of the added carbon was respired, resulting in a highly enriched  ${}^{13}C_{DIC}$  pool, enabling the tracing 147 of autotrophic processes in addition to heterotrophic processes. 148

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Figure 1. Arabian Sea depth gradients. Shown are  $\delta^{13}C_{org}$  and % TOC values of core top 152

sediments, and bottom water oxygenation plotted with depth (a-d), all but  $\delta^{13}C_{org}$  replotted from 153 Lengger et al. (2014), a scatter plot of oxygen exposure time versus  $\delta^{13}C_{org}$  in Arabian Sea core

154 tops along Murray Ridge (e), and a map of the sampling stations with the two main stations used

155

for BHT analysis used here indicated with a star (f). 156

Eight cores are discussed here; these were incubated under oxic or suboxic conditions for 157 7 days (125  $\mu$ M, 6  $\mu$ M O<sub>2</sub>, respectively; Table 1). At the end of incubation, cores were sliced in 158 the intervals 0 - 2, 2 - 4, and 4 - 10 cm depth. They were then frozen and freeze dried for the 159 isotopic analysis of the bacteriohopanepolyol lipids (BHPs), including bacteriohopanetetrol 160 (BHT) and its stereoisomer and biomarker for anammox bacteria, BHT'. Both biomarkers have 161 been studied previously in the Arabian Sea water column and sediments (Jaeschke et al., 2009; 162 Sáenz et al., 2011) Furthermore, cores from 8 stations between 900 and 3000 m water depth were 163 also collected, as described by Lengger et al. (2014, 2012b). The top 0 - 0.5 cm were used for 164 total organic carbon (TOC) and <sup>13</sup>C of organic carbon analysis. For the core from P900 (32 cm 165

length), all depths were analysed in 0.5 cm - 4 cm resolution (Lengger et al., 2012b).

167 2.2 Anammox enrichment cultures

To determine the  $\delta^{13}$ C values for the two bacteriohopanetetrols in anammox bacteria (BHT and BHT', the latter being unique to anammox in the marine environment), an enrichment culture of '*Ca*. Scalindua profunda' was analysed. It was grown in a sequencing batch reactor as described by van de Vossenberg et al. (2008). Analysis of this enrichment culture that showed '*Ca*. S. profunda' comprised about 80% of the cells, while other bacteria belonging to the phyla Bacteriodetes (including Flavobacteriaceae) and Proteobacteria (including Alphaproteobacteria) accounted for the majority of the remaining populations (van de Vossenberg et al., 2008, 2013).

175 2.3. Extraction and purification

The freeze-dried subsamples of the unamended and incubated cores were ground, and the 176 homogenised sediments and the culture were extracted by a modified Bligh-Dyer extraction 177 method (Lengger et al., 2012a). Briefly, they were extracted ultrasonically three times in a 178 mixture of methanol/dichloromethane (DCM)/phosphate buffer (2:1:0.8, v:v:v) and centrifuged, 179 and the solvent phases were combined. The solvent ratio was then adjusted to 1:1:0.9, v:v:v to 180 separate the DCM phase. Liquid extraction was repeated two more times, the DCM fractions 181 were combined, the solvent was evaporated and the larger particles were filtered out over glass 182 wool. The extraction procedure was performed on the enrichment culture material and repeated 183 on the sediment for analysis of BHPs. An aliquot of the extract was subjected to column 184 chromatography using 5% aminopropyl solid phase extraction (SPE), eluting with hexane, DCM, 185 and MeOH, which contained BHPs. For analysis by chromatographic techniques, the extract was 186 derivatised in 0.5 mL of a 1:1 (v:v) mixture of acetic anhydride and pyridine at 50 °C for 1 h, 187 then at room temperature overnight in the case of HPLC-MS analysis. Solvent was dried under a 188 stream of N<sub>2</sub> on a 50°C heating block. 189

190 2.4. Instrumental techniques

191 2.4.1. High temperature gas chromatography coupled to flame ionization detection (HTGC-FID)

GC analysis of acetylated BHPs was done using a HP-5890 Series II GC equipped with a flame ionization detector was fitted with a 0.25 mm x 0.1  $\mu$ m VF5-ht capillary column (CP9045, CP9046, Agilent Technologies UK Ltd., Stockport, UK) of 30 m length (Lengger et al., 2018). An on-column injector was used. To the 30 m column, 1m of a 0.25 mm HT-deactivated silica tubing was attached as a guard column (Zebron Z-Guard, 7CG-G000-00GH0, Phenomenex, Macclesfield, UK). Analysis of bacteriohopanepolyols employed a constant flow of 2 ml/min He and a temperature ramp from 70°C (1 min hold) to 400°C at 7°C min<sup>-1</sup> (1 min hold). 2.4.2. High temperature gas chromatography coupled to mass spectrometric detection (HTGC-MS)

Analysis of acetylated BHPs, using HTGC-MS was performed using a Thermo Scientific 201 Trace 1300 gas chromatograph coupled with an ISO single quadrupole mass spectrometer. 202 Diluted samples were introduced using a PTV injector in splitless mode onto a 0.53 mm fused 203 silica pre-column connected to a 30 m  $\times$  0.25 mm i.d. fused-silica capillary column coated with 204 dimethyl polysiloxane stationary phase (ZB-5HT; film thickness, 0.1 µm; 7HG-G015-02, 205 Phenomenex, Macclesfield, UK). The initial injection port temperature was 70 °C with an 206 evaporation phase of 0.05 min, followed by a transfer phase from 70 °C to 400 °C at 0.2 °C s<sup>-1</sup>. 207 The oven temperature was held isothermally for 1 min at 70 °C, increased at a rate of 7 °C min<sup>-1</sup> 208 to 400 °C and held at 400 °C for 10 min. Helium was used as a carrier gas and maintained at a 209 constant flow of 2.5 ml min<sup>-1</sup>. The mass spectrometer was operated in the electron ionization 210 (EI) mode (70 eV) with a GC interface temperature of 400 °C and a source temperature of 211 340 °C. The emission current was 50 µA and the mass spectrometry set to acquire in the range of 212 m/z 50–950 Daltons with 0.5 s dwell time. Data acquisition and processing were carried out 213 using the Thermo XCalibur software (version 3.0.63). Due to the lack of authentic standards for 214 BHT and BHT', only relative and not absolute values are reported, assuming similar ionization 215 energies. 216

2.4.3. High temperature gas chromatography coupled to isotope ratio mass spectrometry (HTGC-218 IRMS)

The stable carbon isotopic composition ( $\delta^{13}$ C) of BHPs were determined using HTGC-219 isotope ratio mass spectrometry. To this end, an Elementar visION IRMS with GC5 interface 220 (Elementar UK Ltd., Cheadle, UK), and an Agilent 7890B GC were modified in-house and 221 allowed us to achieve column temperatures of up to 400 °C, which resulted in baseline resolution 222 of BHT and BHT' (Fig. 2). 1 µl of the derivatized samples dissolved in ethyl acetate were 223 injected on a cool-on-column injector, into a Zebron Z-Guard Hi-Temp Guard Column (1 m x 224 0.25 mm, Zebron Z-Guard, 7CG-G000-00GH0, Phenomenex, Macclesfield, UK) and separated 225 on a Zebron ZB-5HT analytical column (30 m x 0.25 mm x 0.1 µm, Phenomenex Ltd., 226 Macclesfield, UK). He was used as a carrier gas at a flow rate of 1.5 ml min<sup>-1</sup> and the oven was 227 programmed as follows: 1 min hold at 70 °C, increase by 7 °C min<sup>-1</sup> to 350 °C (10 min hold). 228 Organic compounds were combusted to  $CO_2$  in a 0.7 mm ID quartz tube with CuO pellets at 229 850°C. Instrumentation performance was monitored using an n-alkane standard (B3, A. 230 Schimmelmann, Indiana University, Bloomington, IN, USA; RMS 0.4 ‰), and results were 231 232 calibrated using an in-house mixture of five fatty acid methyl esters, which was injected between every six sample analyses and analyzed using a He flow of 1 ml min<sup>-1</sup>, with a slightly different 233 temperature program (injection at 50 °C held for 1 min followed by an increase of 10°C min<sup>-1</sup> to 234 235 300 °C and a 10 min hold). This is the first time baseline resolution between BHT and BHT' has been achieved on a GC-IRMS, which allows the direct determination of the isotopic composition 236 of both BHT and BHT' in sediment samples (Fig. 2). The isotopic composition of the acetyl 237 group used to derivatise the BHT and BHT' was determined by acetylation of *myo*-inositol, and 238 then subtracted from the values of BHT and BHT' in a mass balance correction (Angelis et al., 239 2012), as authentic standards for BHT or BHT' were not available. 240



#### 242

Figure 2. HTGC-IRMS chromatogram showing baseline separation between BHT, BHT', and
 BHpentol.

245 2.4.4. High performance liquid chromatography coupled to positive ion atmospheric pressure246 chemical ionization mass spectrometry (HPLC/APCI-MS)

To verify the GC-derived assignments, an aliquot of the acetylated BHP samples was dissolved in MeOH:propan-2-ol (3:2; v:v) and filtered on 0.2  $\mu$ m PTFE filters. BHPs were analysed by HPLC/APCI-MS, using a data-dependent scan mode (3 events) on an HPLC system equipped with an ion trap MS, as described in Talbot et al. (2007) and van Winden et al. (2012). Relative BHP concentrations were semi-quantitatively estimated based on the response factor of authentic standards (M. Rohmer; Strasbourg, France; Cooke et al., 2008), with a typical

reproducibility of  $\pm 20\%$ , according to Cooke et al. (2009).

254 2.4.5. Bulk sedimentary organic matter and suspended particulate matter

Freeze-dried core tops from 8 stations between 900 and 3000 m for sedimentary organic matter, and punches from 0.7  $\mu$ m GFF filters for suspended particulate organic matter, were decalcified with 2N HCl, washed, freeze-dried, and subjected to analysis via a Flash EA 1112 Series (Thermo Scientific) analyser, coupled via a Conflo II interface to a Finnigan Delta<sup>plus</sup> mass spectrometer as described by Lengger et al. (2014; sediment) and Pitcher et al. (2011; filters). Standards for  $\delta^{13}$ C analysis were acetanilide and benzoic acid and samples were analysed in duplicate.

#### 3. Results 262

To quantify the provenance of sedimentary organic matter and the contribution of 263 anammox from the OMZ, we analysed the isotopic composition of sedimentary organic matter 264 deposited in close spatial proximity over a large depth gradient in the Arabian Sea, as well as the 265 isotopic composition of biomarker lipids derived from anammox bacteria, chemoautotrophic 266 microbes living in the OMZ. In order to determine whether these were water-column derived or 267 sedimentary, we used stable isotope probing experiments on sediments retrieved from within and 268 below the OMZ. 269

- $\delta^{13}$ C values of C<sub>org</sub> 3.1. 270
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 $\delta^{13}$ C values of C<sub>org</sub> in surface sediments were low (-21.5 ‰) at P900 and increased with 272 water depth to -20.2 ‰ at P2500 (Fig. 1d).  $\delta^{13}C_{org}$  values correlated positively and linearly 273 (Slope 0.0057,  $R^2 = 0.90$ , Figure 1e) with oxygen exposure times as calculated by Koho et al. 274 (2013) and Lengger et al. (2014). Similarly, organic carbon content in the core tops was 275 negatively correlated with oxygen exposure times ( $R^2 = 0.93$ , from Lengger et al., 2014). The 276 increase mirrors the decrease in % TOC – and thus progressing degradation – with increasing 277 oxygen exposure time (Fig. 1b,c, Lengger et al., 2014). At P900, where the whole depth of the 278 core was analysed, values increased slightly with depth, from -21.5 ‰ at the surface, to -20.9 ‰ 279 (Fig. 3).  $\delta^{13}$ C values of particulate organic carbon (suspended particulate organic matter) 280 decreased throughout the water column from -19 to -21.8 %, though with a substantially  $^{13}$ C-281 depleted, yet unexplained, value at the very surface (20 m depth) of -22.9 ‰ (Fig. S1). 282 283



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Figure 3. Depth profile of  $\delta^{13}C_{org}$  and % TOC in an unamended core at P900. 285

- 286 3.2. **Biomarkers** 287
- 288

289 3.2.1. Bacteriohopanepolyols (BHPs)

Analysis of BHPs in the Arabian Sea cores using both HPLC-APCI-MS and HTGC-MS, 290 all showed that BHT', specific for Scalindua, was abundant, and the fractional abundance of 291 BHT's when compared to the sum of BHT and BHT' ranged from 0.4 to 0.6. Other, relatively 292 non-source-specific, BHPs were also present: BHT, 35-aminobacteriohopane-32,33,34-triol 293

294 (aminotriol), bacteriohopane-31,32,33,34,35-pentol (BHpentol), bacteriohopanetetrol cyclitol

ether (BHT-CE) and anhydro-BHT (Fig. S3). The relative concentrations of

bacteriohopanetetrols were 75 to 96 % of total bacteriohopanepolyols, an order of magnitude

higher than other BHPs (Fig. S3). HTGC-MS was able to detect BHT, BHT' and BHpentol (Fig.

S2), as well as small amounts of anhydroBHT, a BHT degradation product, and BHP-570, which

has been tentatively identified by Sessions et al. (2013) as acetylated bacteriohopanediol,

possibly also a degradation product of bacteriohopanetetrol. In the core from P900, the ratio of DUT: even DUT in proceed with doubt (Fig. S4)

301 BHT' over BHT increased with depth (Fig. S4).

We also analysed the BHP content of sediment cores incubated with <sup>13</sup>C-labelled organic matter at both sites under the different conditions detailed by Pozzato et al. (2013 a, b), which was respired and generated <sup>13</sup>C-labelled DIC, allowing to trace autotrophic processes such as anammox. No changes were noted, except for in P900; in those, BHT abundance increased (i.e. BHT'/BHT decreased), indicating that some of the sedimentary BHT could have been produced in situ (Fig. S4).

308 3.2.2. BHT and BHT'  $\delta^{13}$ C values

To establish the isotopic difference of BHT and BHT' derived from anammox we 309 analysed biomass obtained from a batch reactor. BHT and BHT' were the main biohopanoids in 310 the biomass from 'Ca. S. profunda' detected by HTGC-MS (Fig. S2b). In addition to being 311 312 present in 'Ca Scalindua sp.' anammox bacteria, BHT is a ubiquitous lipid common to many bacteria, BHT', however, is specific to Ca Scalindua sp.' in marine environments (Rush et al., 313 2014). The  $\delta^{13}$ C values of BHT and BHT' were identical within the error of analysis (-49 and -314 48 ‰, respectively; Table 1), indicating identical fractionation and thus biosynthetic pathways 315 for both lipids. 316

In the Arabian Sea sediments (all unamended cores), BHT was markedly enriched in <sup>13</sup>C 317 relative to BHT', with values ranging from -24.7 to -28.8 ‰ and -39.1 to -48.1 ‰, respectively 318 (Figs. 4b-c). At P1800 (below the OMZ), BHT and BHT' were slightly more depleted in <sup>13</sup>C, 319 with BHT at  $-27 \pm 3$  ‰ and BHT' at  $-47 \pm 4$  ‰, as compared to  $-26 \pm 1$  ‰ and  $-43 \pm 5$  ‰ for 320 BHT and BHT' at P900 (in the OMZ). However, the difference between P900 and P1800 was 321 not statistically significant for either BHT or BHT'. Moreover, even though the proportion of 322 BHT' increased with depth in the anoxic core, the  $\delta^{13}$ C values did not change. We also analysed 323 BHT and BHT' in the cores that had been incubated with <sup>13</sup>C-labeled POM and DOM, and these 324 showed no indication of <sup>13</sup>C-enrichment in BHT'. Excluding outliers (defined by a Grubbs test at 325 99% confidence level and indicated in Table 1), BHT' values were on average  $-48 \pm 4$  ‰ and -326  $46 \pm 2$  ‰ at P1800, and P900, respectively. BHT was slightly enriched compared to the 327 unamended incubations at 3 cm depth ( $\Delta \delta^{13}$ C = 4.6 ± 0.7 ‰), but not at 1 cm depth. BHpentol 328 concentrations were too low to allow reliable isotopic determination, and anhydroBHT and BHP-329 570 co-eluted with other compounds, also precluding their isotopic characterization. 330

### **331 4. Discussion**

4.1. Origins of BHT and BHT' - a biomarker for anammox

BHT', a biomarker likely unique for annamox in marine environments (Rush et al., bight 2014,), is highly abundant in the sediment and occurred throughout both sediment cores (Fig. 4a), suggesting a significant contribution of anammox biomass to sedimentary organic matter. In

- the anoxic core (P900), BHT and BHT' were present in concentrations at least an order of
- magnitude higher than other BHPs (Fig. S2), and BHT' was the most abundant of the two
- stereoisomers (Fig. 4a). The high fractional abundance of BHT' over the sum of BHT and BHT' (0.4 - 0.6) is contrary to our expectations, as BHT is a ubiquitous lipid and presumed to derive
- from both anammox and non-anammox sources such as cyano- and many other bacteria (Pearson
- and Rusch, 2009); it is, therefore, expected to be abundant in most depositional contexts. Lower
- proportions of BHT were reported previously in nearby core tops (0.22-0.30) by Saenz et al.
- 343 (2011); this could be caused by a difference in settings, or in BHP extraction protocol. However,
- the high proportions of BHT' observed in the Arabian Sea are not unprecedented: they are
- slightly lower than BHT' proportions in sediments underlying the Humboldt Current System
- OMZ offshore Peru (0.45 0.69 in surface sediments, Matys et al., 2017).



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> Figure 4. Anammox biomarkers in unamended Arabian Sea sediment. Panel (a) shows the proportion of BHT' relative to BHT+BHT', panels (b) and (c) show the  $\delta^{13}$ C values of BHT and BHT' in the unamended oxic (P1800) and anoxic core (P900), respectively.

352

Supporting evidence for the unique source of BHT' comes from its <sup>13</sup>C values of -40 to -50‰. Anammox bacteria are known to fractionate strongly against  $\delta^{13}$ C, with up to 26 ‰ fractionation observed for biomass in cultures and sediment, with even more strongly depleted

BHT and BHT' (Brocadia sp. by 47 ‰ and Scalindua sp. by 49 ‰ against DIC; Schouten et al.,

357 2004). Anammox bacteria within the oxygen minimum zone and using dissolved inorganic

carbon, which is up to 2 ‰ lighter within the OMZ than at the surface (Kroopnick, 1984), are
likely to produce such <sup>13</sup>C-depleted lipids. In line with an anammox origin, BHT' here is
decidedly more depleted than other biomarkers in the Arabian Sea (cf. Wakeham and McNichol,
2014), though distinctly depleted highly branched isoprenoids (HBIs, -37 ‰) have been found in
Arabian Sea cores from the Holocene (Schouten et al., 2000).

BHT, however, is much more enriched in <sup>13</sup>C in the Arabian Sea sediments, with values 363 of -29 to -30 ‰ (Fig. 4b,c). This is in contrast to anammox cultures, where BHT and BHT' are 364 produced with identical isotope values (Fig. S4, Table 1). This strongly supports the idea that 365 BHT has a mixed origin. The sources for BHT could be varied: cyanobacteria or heterotrophic 366 bacteria thriving in the OMZ or the sediment (Pearson and Rusch, 2009), and possibly including 367 nitrite-oxidising bacteria (Kharbush et al., 2018). The  $\delta^{13}$ C values of BHT derived from these 368 alternative sources are poorly constrained (Hayes, 2001; Kharbush et al., 2018; Pearson, 2010; 369 Sakata et al., 1997; Schouten et al., 1998). However, for heterotrophic bacteria, values similar to 370 the consumed OM minus the depletion associated with polyisoprenoids of 6-8 ‰ are expected 371 (Pancost and Sinninghe Damsté, 2003). Similar values would be expected for cyanobacterial 372 lipids, too, as those are depleted by 22 - 30 % compared to dissolved CO<sub>2</sub>, which varies from 0 373 to 2 ‰ in the photic zone. Methane cycling could also lead to <sup>13</sup>C depleted OM, but is not 374 important in the Arabian Sea (Lüke et al., 2016), and there is no biomarker evidence for it, such 375 as <sup>13</sup>C depleted archaeal lipids, or methylated BHPs. Thus, a BHT  $\delta^{13}$ C value of -30 % suggests 376 heterotrophic and other bacterial sources, possibly mixed with an anammox source. 377

378 Anammox activity has been shown to occur both within the OMZ water column (Jensen et al., 2011; Lüke et al., 2016; Pitcher et al., 2011; Villanueva et al., 2014) as well as in 379 sediments (Sokoll et al., 2012; Devol, 2015). To test whether the anammox-derived biomarkers 380 are formed in the sediment, the labelled cores were incubated for 7 days with particulate and 381 dissolved organic matter (Table 1), which resulted in the substantial incorporation of <sup>13</sup>C into 382 bacterial fatty acids, as well as generating  ${}^{13}$ C-enriched CO<sub>2</sub> due to heterotrophic respiration (up 383 384 to 14 % of the added C was respired, Pozzato et al., 2013a,b). Anammox bacteria are autotrophic, and it is therefore likely that an active sedimentary community would result in some 385 uptake of this <sup>13</sup>C labelled CO<sub>2</sub> formed by respiration. However, no significant labelling was 386 observed in BHT' or BHT (Table 1, Table S1), suggesting that most of this pool is water-column 387 derived. Further, BHT and BHT' are also present in the surface of the oxic sediments (P1800), in 388 proportions similar to the anoxic sediments at P900. Even if anammox growth was too slow for 389 labelling to take effect, substantial sedimentary production would have resulted in a decreasing 390  $\delta^{13}$ C value of BHT' in the unamended cores, as DIC gradually becomes more <sup>13</sup>C depleted with 391 sediment depth (Fernandes et al., 2018), but this is not observed (Fig. 4bc). Collectively, these 392 data support the idea that the vast majority of BHT' is derived from anammox bacteria living in 393 the OMZ of the water column, and that BHT also has a predominant pelagic origin with limited 394 sedimentary production. 395

The  $\delta^{13}$ C values of geohopanoids (i.e. the geological degradation product of biohopanoids such as BHT and BHT') can exhibit dramatic variability, and often pronounced depletion in terrestrial (Pancost et al., 2007) and marine settings (Köster et al., 1998). These are commonly attributed to aerobic methane oxidising bacteria, and thus regarded as evidence for a significant contribution of methane oxidisers to sedimentary organic matter. However, our data show that hopanes of -35 to -50 ‰ could also indicate the presence of a significant amount of anammox bacteria in an anoxic water column. Several factors could attenuate the anammox signal. The decrease of

biohopanoids in structural complexity upon degradation means that the anammox signal would 403 be diluted by mixing with aerobically and anaerobically produced hopanoids such as those of 404 Geobacter (Fischer et al., 2005; Härtner et al., 2005). Chemoautotrophs operating in euxinic 405 settings employ biochemical pathways resulting in <sup>13</sup>C-enriched biomass and lipids compared to 406 DIC (van Breugel et al., 2005b). However, in anoxic basins such as the Black Sea or anoxic 407 fjords, remineralization of organic matter also results in distinctly depleted  $\delta^{13}C_{DIC}$  below the 408 chemocline (-12 ‰; Fry et al., 1991; Volkov, 2000). This may explain why, in some marine 409 anoxic settings where we might expect to see an anammox signature, the <sup>13</sup>C depletion of 410 hopanes parallels that of algal biomarkers (Sinninghe Damsté et al., 2008; van Breugel et al., 411 2005a). Nonetheless, we suggest that potential anammox contributions to the sedimentary 412 413 hopanoid pool should be considered when interpreting their abundances, distributions and

414 isotopic compositions.

#### 415 4.2. Origin of sedimentary organic matter

The Murray Ridge represents an open ocean setting, and the selected coring sites were in 416 close proximity to each other, with no substantial terrigenous contribution (Koho et al., 2013; 417 Lengger et al., 2014, 2012b; Nierop et al., 2017; Fig. 1d). Despite this,  $\delta^{13}$ C values of 418 sedimentary organic matter of a purely marine origin varied: The value at the shallowest location 419 (-21.5 ‰; P900), within the OMZ, was 2.5 ‰ more depleted than the estimated value for surface 420 water-derived OM -19.8 ‰ (Fontugne and Duplessy, 1986), and the  $\delta^{13}$ C values of sedimentary 421 Corg increased with increasing oxygen bottom water concentrations / oxygen exposure time (Fig. 422 1e). These observations agree with earlier studies from this setting. Cowie et al. (2009, 1999) 423 detected similar trends in surficial sediments across different settings in the Arabian Sea, with 424 values of -21 ‰ within and -19 ‰ above and below the OMZ. Organic matter in sediment traps 425 collected in the north western Arabian Sea (i.e. sinking POC) had a  $\delta^{13}$ C value of -22.4 ‰ 426 (composite of the OMZ between 500 and 900 m depth). The corresponding sedimentary  $\delta^{13}C_{org}$ 427 value, from oxygenated bottom waters at 1445 m depth, was, however, more enriched (-20.8%); 428 Wakeham and McNichol, 2014). Fernandes et al. (2018) detected similar, though less 429 pronounced, trends in sediments collected from the Pakistan margin. An increase in  $\delta^{13}$ C with 430 enhanced degradation, i.e. <sup>13</sup>C-enriched sediments vs. depleted POM, was also observed in the 431 South China Sea (Liu et al., 2007), and in the Eastern Tropical North Pacific (Jeffrey et al., 432 1983). 433

Despite its common occurrence in OMZ settings, this trend is unusual and, at present, not 434 explained: degradation of organic carbon in marine environments usually preferentially removes 435 isotopically heavy carbon (Hatch and Leventhal, 1997), causing a depletion in  $\delta^{13}$ C with 436 increased degradation of the sediment. This can be due to the removal of the more labile marine 437 carbon, and subsequent relative enrichment of terrigenous organic material of a lower initial 438 reactivity and lower  $\delta^{13}$ C values (Huguet et al., 2008; Middelburg et al., 1993). However, 439 progressive depletion also occurs in areas with purely marine input; this is due to preferential 440 loss of <sup>13</sup>C-enriched carbohydrates over the more <sup>13</sup>C-depleted lipids, and preferential 441 degradation of easily accessible material over biopolymers (Spiker and Hatcher, 1987), and 442 polymerization and elimination of functional groups (Galimov, 1988; Balabane et al., 1987). 443 Conversely, sulfurization, a process observed in euxinic settings appears to preferentially 444 preserve <sup>13</sup>C-enriched material such as carbohydrates (Van Kaam-Peters et al., 1998); however, 445 this process is not expected to occur here - the Arabian Sea is anoxic but not sulfidic (Ulloa et 446 al., 2012) and has not experienced euxinia for the past 120 ka (Schenau et al., 2002). 447

Nonetheless, in Arabian Sea sediment,  $\delta^{13}C_{org}$  values increased by 1.8 ‰ with increasing oxygen 448 exposure time and thus increasing degradation; at the same time, organic carbon contents 449 decreased from 60 to 10 mg g dw<sup>-1</sup>, indicating progressing remineralization (Fig. 1c, Lengger et 450 al., 2014). Cowie (2005) attributed this to the contribution of – potentially – organic matter from 451 the facultatively autotrophic, chemosynthetic sulfur-bacterium Thioploca sp., which has been 452 observed in the Arabian Sea (Schmaljohann et al., 2001). However, Thioploca sp. has only been 453 reported for shelf and upper slope sediments in the Arabian Sea (above and upper part of OMZ). 454 and it is unlikely that the sulfur-dependent Thioploca sp. could have caused this significant 455 depletion by chemoautotrophy, as sulfide concentrations are negligible within the OMZ (Kraal et 456 al., 2012), and there is no evidence for the production of severely <sup>13</sup>C-depleted biomass by 457 filamentous sulfur bacteria (Zhang et al., 2005). Further, the depletion is not only observed in 458 sediments, but also in particle fluxes (Wakeham and McNichol, 2014), which strongly suggests 459 the exclusion of sedimentary sources for <sup>13</sup>C-depleted organic matter. This is also supported by 460 the  $\delta^{13}C_{org}$  values of the suspended particulate matter (Fig. S1), which becomes gradually more 461 depleted with depth. 462

An active nitrogen-cycle in the water column within the Arabian Sea OMZ is 463 undisputedly present, with heterotrophic denitrifying bacteria, nitrifying archaea, nitrite-464 oxidising and anammox bacteria present in high abundances (Lüke et al., 2016; Villanueva et al., 465 2014). The  $^{13}$ C fractionation for the carbon fixation pathways employed by nitrifying archaea (3-466 Hydroxypropionate/4-Hydroxybutyrate, 3-HP/4-HB) is similar to that of phytoplankton using 467 Rubisco (Könneke et al., 2012). The biomass of heterotrophic denitrifiers is close to the value of 468 the source organic matter (1 ‰ more enriched; Hayes, 2001), and the biochemical pathway 469 employed by nitrite oxidisers (reverse TCA cycle) produces <sup>13</sup>C-enriched biomass (Pearson, 470 2010). Anammox bacteria, however, are abundant and active in the Arabian Sea (Jensen et al., 471 2011) and known to produce highly <sup>13</sup>C-depleted biomass by inorganic carbon fixation 472 (Schouten et al., 2004). Anammox bacteria, and other, yet undescribed chemoautotrophs, could 473 thus present a potential pathway for addition of <sup>13</sup>C-depleted organic matter to sinking organic 474 matter, a hypothesis we explore further below. 475

The water-column derived <sup>13</sup>C-depleted BHT', in combination with the unusual  $\delta^{13}C_{org}$ 476 trends, suggests that there may be a substantial contribution of <sup>13</sup>C depleted organic carbon, 477 produced by anammox bacteria, to the sedimentary organic matter. Based on BHT'  $\delta^{13}$ C values 478 determined in this study, and the fractionation factor associated with anammox lipid biosynthesis 479 of 16 % (lipid versus total biomass; Schouten et al., 2004), we can estimate a  $\delta^{13}$ C value for 480 anammox biomass: of ca.-28.6  $\pm$  6 ‰, which is similar to the expected value calculated from 481 ε<sub>biomass-DIC</sub> (22-26 ‰; Schouten et al., 2004) and the generally observed DIC value in the Arabian 482 Sea OMZ at depth, 0 ‰ (Moos, 2000). This is significantly depleted compared to phytoplankton 483 biomass in the Arabian Sea (-19.8 ‰, Fontugne and Duplessly, 1978), and would add depleted 484 organic carbon to the sinking POM. 485

We modelled the contribution of annamox biomass ( $\chi_{anammox}$ ) to SOM using an isotopic mass balance approach (using IsoError; <u>Phillips et al., 2005</u>), which employs uncertainty propagation and error estimates and allows the determination of contributions of likely sources of organic matter to the sediment. The end member value of  $-19.8 \pm 0.5$  ‰ was used for phytoplankton-derived SOM (Fontugne and Duplessy, 1986; Ziegler et al., 2008). This value is slightly lower than the -19 ‰ observed from the surface particulate carbon, but is representative of sedimentary organic matter, as it accounts for the decreased  $\delta^{13}$ C values of POM caused by degradation of <sup>13</sup>C enriched compounds such as carbohydrates (Close, 2019; Ziegler et al., 2008). More recent values for phytoplankton  $\delta^{13}$ C are, to the best of our knowledge, not

495 available. Planktonic, sinking organic matter, and organic material produced by heterotrophs also

contribute to sedimentary OM. However, this OM value would likely be similar to the organic

matter assimilated (Blair et al., 1985; 2001).  $-28.6 \pm 6$  ‰ was used for anammox-derived organic

498 carbon, as derived from the  $\delta^{13}$ C value of BHT' and  $\varepsilon_{bm/lipid}$  of  $16 \pm 4$  % (Schouten et al., 2004; 499 Table 1; uncertainty represents combined standard deviations of  $\delta^{13}C_{BHT'}$  and  $\varepsilon_{lipid/bm}$ ). The  $\delta^{13}C$ 

value -21.5% of surface sediment  $C_{org}$  in the OMZ was used as the value of the mixture (Table 1):

502

 $\chi_{anammox} \cdot \delta^{13} C_{anammox} + \chi_{PP} \cdot \delta^{13} C_{PP} = -21.5 \%$ 

Ean 1

The modelling, assuming the above end member contributions and their statistical uncertainties, 505 yields a proportion of anammox with a mean of approximately 17% (Fig. 5), and confirms that, 506 with 95% confidence, anammox contribution to the sedimentary organic matter is between 3 and 507 30% among the different cores. This suggests that some of the sedimentary organic matter 508 (SOM) present at P900 is anammox-derived. However, other chemoautotrophic bacteria also 509 present, or suspected to be present, in the Arabian Sea OMZ (e.g. ammonia-oxidizing archaea, 510 Pitcher et al., 2011) possess metabolisms which would lead to different  $\delta^{13}$ C values, and could 511 therefore be diluting this signal (Hayes, 2001; Pearson, 2010). These results suggest that the 512 contribution of prokaryotic organic material produced in the OMZ to SOM is larger than 513 estimated in this simple, two-component mass balance. A three-source model resolving 514 heterotrophic bacteria or degraded OM, anammox bacteria and phytoplankton was solved with 515 SIAR (Parnell et al., 2010) and yielded similar results (not shown). 516



**Figure 5.** Results from the isotope mass balance model, showing the calculated means, standard error, and confidence intervals for contribution from anammox, and planktonic-derived OM

520 (POM) to the surface sediment at P900.

521

This estimate, at a first glance, appears large. However, we can convert experimental 522 rates of anammox-mediated ammonia oxidation of 27 - 38 nmol L<sup>-1</sup> d<sup>-1</sup> as determined in the 523 Arabian Sea (Jensen et al., 2011) and alternative rates of 0.24 -4.32 nmol L<sup>-1</sup> d<sup>-1</sup> estimated by 524 Ward et al. (2009), to carbon fixed using stoichiometric rates determined for anammox bacteria 525 of 0.07 mol C mol  $N_2^{-1}$  (Jetten et al., 2001; Strous et al., 1999; van de Vossenberg et al., 2008). 526 Assuming a production depth of 300 to 900 m water depth, with a maximum at 600 m (Pitcher et 527 al., 2011), anammox production over the whole depth could be estimated using Equation 2 (see 528 529 Fig. S5),

530

$$P = \frac{(k \cdot 600)}{2}$$

532

in which *k* corresponds to maximum rate at maximum production depth calculated from aforementioned published anammox rates (*k* equals  $138.7 - 9855 \mu mol m^{-3} yr^{-1}$ , Jensen et al. 2011;  $87.6 - 1576.8 \mu mol m^{-3} yr^{-1}$ , Ward et al. 2009; Fig. S5).

Eqn. 2

Using this equation, we calculated that water-column anammox bacteria produce up to 536 3.5 g organic C m<sup>-2</sup> yr<sup>-1</sup>. These estimates dwarf sedimentary anammox rates, with C-fixation 537 determined to be at 64 pg organic C m<sup>-2</sup> yr<sup>-1</sup> in the Arabian Sea (Sokoll et al., 2012), indicating 538 that most of the anammox carbon in the sediment is water-column derived. Given organic carbon 539 accumulation rates (Lengger et al., 2012b) of 3 to 5 g C m<sup>-2</sup> yr<sup>-1</sup> at P900 and assuming a 17 % 540 anammox contribution to sedimentary organic matter, a maximum of 24 % of annually produced. 541 water-column anammox biomass is preserved in this anoxic setting. Within the sediment from 542 the OMZ, as seen from a sedimentary depth profile at P900 in the OMZ (Fig. 3),  $\delta^{13}C_{org}$  also 543 shifts to higher values with depth, to a maximum of -20.8 ‰ at 8 cmbsf. However, contrary to 544 oxic degradation, no decrease in SOM contents occurs over the first 14 cm (Fig. 3). In this case, 545 the increase in  $\delta^{13}C_{org}$  likely reflects input by other sedimentary autotrophs, producing enriched 546 organic carbon. This suggests that the <sup>13</sup>C-depleted SOM signal from the OMZ is not completely 547 preserved. 548

549 Other chemoautotrophs, where specific biomarkers are not available, could also contribute to the SOM: at depth, the increased availability of CO<sub>2</sub> derived from heterotrophic 550 respiration of the POM pool would result in greater discrimination against <sup>13</sup>C (Freeman et al., 551 1994; Freeman, 2001). An example are lipids of ammonia-oxidising archaea, which were more 552 depleted when deposited in the OMZ than below the OMZ (Table S2), suggesting that some of 553 those were produced in the produced in the OMZ and transported to the sediment (Lengger et al., 554 2014, 2012b; Schouten et al., 2012).. However, as these archaeal lipids present a mixed pelagic 555 and sedimentary signal, quantitative estimates are not possible. 556

557 Hence, chemoautotrophic carbon fixation could explain the low  $\delta^{13}$ C values of 558 sedimentary OM observed within the OMZ. As the newly produced organic carbon is more 559 labile than surface-produced organic matter, it would degrade more quickly upon exposure to

oxic bottom waters. This poorer preservation of anammox biomass results in a shift back towards 560 the <sup>13</sup>C signature of the primary photosynthetic production, which is observed as the enrichment 561 in sedimentary  $\delta^{13}C_{org}$  with increasing oxygen exposure times (Fig. 1e). A likely explanation for 562 this lability of anammox biomass is that, due to the lack of zooplankton in the OMZ, this organic 563 matter is not fecal-pellet packaged or adsorbed to inorganic particles and thus not matrix 564 protected (cf. Burdige, 2007). Wakeham et al. (2002) analysed the biomarker fluxes in sinking 565 particles in the Arabian Sea, and they also found that surface-produced lipids such as alkenones 566 (which would be fecal pellet packaged) were exceptionally well-preserved compared to the total 567 organic carbon of the SPM. Gong and Hollander (1997) also noted an enhanced contribution of 568 bacterial biomass to sediment deposited under anoxic conditions in the Santa Monica Basin, 569 when compared to sediment located in nearby oxygenated bottom waters. Studies examining 570 carbon fluxes in the Arabian Sea (Keil et al., 2016) and the Cariaco basin (Taylor et al., 2001) 571 invoked the addition of chemoautotrophically derived carbon from a midwater source in order to 572 explain the enhanced carbon fluxes observed, even if oxygen depletion and other factors such as 573 the lack of zooplankton were taken into account. 574

575

#### 576 4.3. Quantifying the biological pump and remineralization rates

577 Models of the biological pump consider primary production, particle fluxes, oxygen

concentrations, and respiratory rates. Sinking organic matter is traditionally regarded as a

reflection of exported organic material from the photic zone, and export models consider

remineralization and C-loss (Cabré et al., 2015; Schlitzer, 2002), based on claims that bacterial

contributions do not sink (Buesseler et al., 2007). However, relying on these parameters,
 estimates of particulate organic carbon flux in the Arabian Sea are generally too low to sustain

estimates of particulate organic carbon flux in the Arabian Sea are generally too low to sustain experimentally observed denitrification, bacterial production, and oxygen deficiency in the

Arabian Sea (Anderson and Ryabchenko, 2009; Andrews et al., 2017; Naqvi and Shailaja, 1993;

Rixen and Ittekkot, 2005). New mechanisms have been discovered that can explain downward

transport of smaller particles and bacteria, such as particle aggregation (Burd and Jackson, 2009),

and mixed layer transport mechanisms (Bol et al., 2018), which enable the consideration of the

contribution of anammox biomass to sinking OM.

589 In previous work, the deficit in modelled and observed C-utilization estimated from

<sup>590</sup> denitrification was as high as 70 g C m<sup>-2</sup> y<sup>-1</sup> (Naqvi and Shailaja, 1993). Including anammox in

models of the nitrogen cycle can account for at least 30% or more of the nitrogen losses, as well

as accounting for organic carbon production, thereby reconciling some of this budget

discrepancy. More recently, the combined inclusion of mesopelagic zooplankton and

heterotrophic bacteria into 3D models resulted in much lower bacterial productivity (4.7 g C m<sup>-2</sup>

 $y^{-1}$ ) than observational estimates of 10.4 g C m<sup>-2</sup>y<sup>-1</sup> (Anderson and Ryabchenko, 2009). Other recent models, combined with experimental observations of particle fluxes, suggest that at some

recent models, combined with experimental observations of particle fluxes, suggest that at some stations, the lower oxycline is associated with 0.4 to 1.8 g C m<sup>-2</sup> y<sup>-1</sup> increases in net carbon fluxes

(Roullier et al., 2014). Horizontal transport, also from nepheloid layers, novel transport

599 mechanisms from the mixed layer, and DOM, have previously been invoked as the missing

600 carbon supply. Here, we estimate that an additional  $3.5 \text{ g m}^{-2} \text{ y}^{-1}$  of organic carbon are produced

by anammox from DIC, with other chemoautotrophic processes possibly adding to this estimate.

This also affects estimates of nitrogen losses and has direct implications for the forecasting of

603 OMZs in a warming world.

In paleoceanography, the  $\delta^{13}$ C values of total organic carbon are commonly used in 604 combination with organic carbon contents in sediment cores to interpret changes in sea surface 605 biogeochemistry and to discern the causes for ocean deoxygenation (Jenkyns, 2010; Meyers, 606 2014). Negative  $\delta^{13}$ C excursions are associated with either atmospheric injection of depleted CO<sub>2</sub> 607 and/or increased CO<sub>2</sub> availability (Pagani, 2005; Pancost and Pagani, 2005), or a higher input of 608 terrigenous organic matter. Positive excursions are inferred to reflect increased burial rates of 609 organic carbon, i.e. the removal of 'light' carbon from the ocean-atmosphere reservoir (e.g. 610 Kuypers et al., 2002; Jenkyns, 1985). An increase in TOC coinciding with a negative carbon 611 isotope excursion has been observed in the sedimentary record numerous times. They are a 612 prominent feature in OAE 1a (Leckie et al., 2002), during the PETM (Zachos et al., 2005), and in 613 the Eastern Mediterranean during Sapropel deposition (Meyers and Arnaboldi, 2008). Our results 614 show that if non-sulfidic OMZs expand to impinge on the sediment, a significant amount of <sup>13</sup>C-615 depleted organic carbon can be preserved, changing the carbon isotope composition of organic 616 matter by up to -1.6 % in the case of Arabian Sea surface sediments, and by -0.5 to -1 % in 617 deeper sediments. When evaluating past events based on bulk  $\delta^{13}$ C values, the incorporation of 618 chemoautotrophically produced carbon must thus be considered. This also further supports 619 claims that calculations of pCO<sub>2</sub> based on differences between  $\delta^{13}$ C of organic and inorganic 620 carbon should not rely on bulk organic carbon that includes bacterial contributions, but instead 621 use compound specific  $\delta^{13}$ C values derived from algae, consistent with previous investigations 622 (Pancost et al., 1999; Pancost et al., 2013; reviewed by Witkowski et al., 2018). By extension, a 623 deviation from the relationship between  $\delta^{13}$ C bulk OM and  $\delta^{13}$ C phytane could be an indicator 624 for the chemoautotrophic contribution to sedimentary OM. 625

Chemoautotrophic metabolism in euxinic water columns, where often very low  $\delta^{13}C_{DIC}$ 626 values are observed (van Breugel et al., 2005a,b; Fry et al., 1991; Volkov, 2000), has previously 627 been invoked as contributing to organic matter in these setting. Anammox, which has been 628 shown to be an important process in the anoxic water column overlying euxinia (Kuypers et al., 629 630 2003), also needs to be considered. Anammox also plays an important role in the nitrogen cycle of Mediterranean sapropel events, even in sapropels where euxinia appears to have occurred 631 (Rush et al., 2019), potentially causing a significant contribution of anammox to the depleted 632  $\delta^{13}C_{org}$  values of these sapropels. Previously, a conceptual model invoking chemoautotrophy has 633 been invoked to explain e.g. the End-Permian (Luo et al., 2014) and the Lomagundi isotope 634 excursion (Bekker et al., 2008). 635

Settings similar to the Arabian Sea, with  $\leq 4.5 \ \mu mol \ kg^{-1} O_2$ , occur in the East Pacific and 636 those settings in total occupy an area of 8.45 to 15 x  $10^{12}$  m<sup>2</sup> (Karstensen et al., 2008; Hattori, 637 1983; Paulmier and Ruiz-Pino, 2009), or a volume of 0.46 x  $10^{15}$  m<sup>3</sup> with 1.148 x  $10^{12}$  m<sup>2</sup> in 638 contact with the sediment (Helly and Levin, 2004). OMZs are expanding in size and in volume 639 (Stramma et al., 2010; Queste et al., 2018), and anammox bacteria play a key role in these 640 settings, particularly because they tolerate higher oxygen concentrations than denitrifiers 641 (Dalsgaard et al., 2014). The labile organic matter added to the sinking carbon can fuel 642 heterotrophic processes, further exacerbating oxygen depletion, and nitrogen loss by 643 denitrification. For accurate future forecasting, it is imperative to reconcile experimental and 644 model-based estimates. Novel, detailed observations explaining the vertical transport of bacterial 645 biomass, and evidence from the isotopic composition of TOC and biomarkers for 646 chemoautotrophic processes, such as shown here for anammox, will enable the further 647 unravelling of the biogeochemical mechanisms underpinning present, past, and future OMZs. 648

#### 649 **5 Conclusions**

Combining the  $\delta^{13}$ C values of BHT', a biomarker from anammox bacteria, and the  $\delta^{13}$ C 650 values of sedimentary organic matter allowed us to estimate the contribution of anammox to 651 SOM in the Arabian Sea. This can be as much as 30 % of organic matter deposited within the 652 OMZ. In the sediments underlying the Arabian Sea,  $\delta^{13}$ C values of total organic matter can shift 653 by 1 to 2 ‰ due to these bacterial contributions, but it remains unclear how well and under what 654 conditions this signature is preserved. Our results suggest that chemoautotrophs (e.g. anammox 655 bacteria) contribute more than previously believed to the burial of carbon in oxygen deficient 656 zones, and remineralization rates are potentially higher than inferred from organic matter 657 decreases. This implies that, when past occurrences of OMZs are evaluated based on  $\delta^{13}$ C values 658 of SOM, a <sup>13</sup>C-depleted contribution of bacteria needs to be considered. Chemoautotrophic 659 carbon fixation thus represents a mechanism of CO<sub>2</sub> removal from the pelagic water column, and 660 contributes to fluxes of sinking organic carbon. It explains some of the mismatches in carbon 661 budgets when experimental and modelling estimates are compared - and it should therefore be 662 included in biogeochemical models predicting feedbacks to a warming world. 663 664 Acknowledgements: SKL was supported by Rubicon fellowship nr. 825.14.014 from the 665 Netherlands Organization for Scientific Research (NWO). MJ, JJM, JSSD, and SS were 666 supported by NESSC OCW/NWO 024 002.001 and MJ, JSSD and DR by SIAM OCW/NWO 667 024 002.002 grants. RSN and DR were supported by NERC grant ANAMMARKS 668 NE/N011112/1 awarded to DR. JB is supported by a NERC GW4+ Doctoral Training 669 Partnership studentship from the Natural Environment Research Council (NE/L002434/1) and is 670 thankful for the support and additional funding from CASE partner, Elementar UK Ltd. We

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- 679
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- 682

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#### 1146

1147	<b>Table 1.</b> $\delta^{13}$ C values of analysed samples. s.d. = standard deviation of three repeat analyses, n.d.
1148	= not determined (single analysis). The values shown are natural abundance from unamended
1149	cores, and from incubations with <sup>13</sup> C labelled DOM and POM, under oxic and suboxic

1150 conditions. \* indicates outliers (Grubbs).

				δ <sup>13</sup> C value [‰ V-PDB]		
Type Sample		BHT	s.d.	BHT'	s.d.	Corg
Biomass	Scalindua sp.	-47.8	1	-47.6	1	
	P900 - 1 cm	-24.7	1.3	-39.1	1.3	-21.5
Arabian	P900 - 3 cm	-27.0	1.4	-48.1	1.1	
Sea	P900 - 7 cm	-26.1	0.4	-41.4	0.5	
Natural	P1800 - 1 cm	-25.1	1.7	-48.1	1.6	-20.3
abundance	P1800 - 3 cm	-28.8	2.1	-46.6	4.1	
	P1800 - 7 cm	-27.1	0.6	-45.5	1.0	
	P900, suboxic, DOM - 1 cm	-27.9	n.d.	-45.9	n.d.	
	P900, suboxic, DOM - 3 cm	-26.6	n.d.	-42.0	n.d.	
	P900, suboxic, POM - 1 cm	-41.6*	n.d.	-75.2*	n.d.	
	P900, suboxic, POM - 3 cm	-26.4	n.d.	-49.8	n.d.	
	P900, oxic, POM - 1 cm	-27.1	n.d.	-51.5	n.d.	
Arabian	P900, oxic, POM - 3 cm	-25.2	n.d.	-47.8	n.d.	
Incubations	P1800, oxic, DOM - 1 cm	-21.3	n.d.	-42.0	n.d.	
	P1800, oxic, DOM - 3 cm	-28.3	n.d.	-45.6	n.d.	
	P1800, suboxic, POM - 1 cm	-52.2*	n.d.	-101.9*	n.d.	
	P1800, suboxic, POM - 3 cm	-28.8	n.d.	-47.1	n.d.	
	P1800, oxic, POM - 1 cm	-30.2	n.d.	-51.3	n.d.	
	P1800, oxic, POM - 3 cm	-28.0	n.d.	-46.1	n.d.	

1151

# **@AGU**PUBLICATIONS

#### Global Biogeochemical Cycles

#### Supporting Information for

### Dark carbon fixation contributes to sedimentary organic carbon in the Arabian Sea oxygen minimum zone

Sabine K. Lengger<sup>1,2,3\*</sup>, Darci Rush<sup>3,5</sup>, Jan Peter Mayser<sup>2</sup>, Jerome Blewett<sup>2</sup>, Rachel Schwartz-Narbonne<sup>4</sup>, Helen M. Talbot<sup>4,†</sup>, Jack J. Middelburg<sup>5</sup>, Mike S.M. Jetten<sup>6</sup>, S. Schouten<sup>3,5</sup>, J. S. Sinninghe Damsté<sup>3,5</sup> and Richard D. Pancost<sup>2</sup>

<sup>1</sup> Biogeochemistry Research Centre, School of Geography, Earth and Environmental Science, University of Plymouth, PL48AA, Plymouth, United Kingdom.

<sup>2</sup> Organic Geochemistry Unit, School of Chemistry, University of Bristol, BS81TS, Bristol, United Kingdom.

<sup>3</sup> NIOZ Royal Netherlands Institute for Sea Research, Dept. of Marine Microbiology and Biogeochemistry, and Utrecht University,1797SZ, Texel, The Netherlands.

<sup>4</sup> School of Natural and Environmental Sciences, Newcastle University, Drummond Building, NE1 7RU, Newcastle-upon-Tyne, United Kingdom.

<sup>5</sup> Department of Earth Sciences, Faculty of Geosciences, Utrecht University, 3508 TA, Utrecht, The Netherlands.

<sup>6</sup> Department of Microbiology, IWWR, Radboud University Nijmegen, 6525 XZ, Nijmegen, The Netherlands

\* corresponding author: Portland Square Bldg, School of Geography, Earth and Environmental Science, University of Plymouth, PL48AA, Plymouth, United Kingdom. Email: sabine.lengger@plymouth.ac.uk, Phone: +44 1752 585966 .

† present address: BioArCh, Environment Building, University of York, YO10 5DD, Heslington, United Kingdom.

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#### Introduction

This supporting information includes additional figures to those presented in the main text. A table showing the  $\delta^{13}$ C values of archaeal lipids is added (Table S1), including information on how this data was acquired (Text S1).



**Figure S1**.  $\delta^{13}C_{org}$  values of organic matter in the Arabian Sea. Sedimentary values (white circles), and suspended particulate matter (black circles) in the water column as sampled by McLane in situ filter pumps are shown.



**Figure S2**. HTGC-MS chromatograms and mass spectra of bacteriohopanepolyols. a Arabian Sea sediment and b '*Ca*. Scalindua profunda' biomass, and mass spectra of identified BHPs: c bacteriohopanetetrol (BHT), d bacteriohopanetetrol stereoisomer (BHT'), and e bacteriohopanepentol (BHpentol), with BHT as the only compound with known stereochemistry.



**Figure S3**. BHP composition in core tops at P1800 and P900 as detected by HPLC-APCIlontrap MS.



**Figure S4**. BHT'/BHT ratios compared between incubated cores and unamended. UA – unamended, OP – oxic incubation, particulate <sup>13</sup>C-labelled OM, OD – oxic incubation, dissolved <sup>13</sup>C-labelled OM, SP – suboxic incubation, particulate <sup>13</sup>C-labelled OM, OD – oxic incubation, dissolved <sup>13</sup>C-labelled OM.



**Figure S5.** Illustration of the estimation of anammox production over depth, with a maximum assumed at 600 m acc. to Pitcher et al. (2011), and no production above 300 and below 900 m water depth.

Δδ <sup>13</sup> C valu	e [‰	V-P	DB]
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	BHT	BHT'	bp-0	bp-2	bp-3
Sta 1, suboxic, DOM - 1 cm	-2.64	-5.59	1.9	-0.211	0.102
Sta 1, suboxic, DOM - 3 cm	0.39	4.98			
Sta 1, suboxic, POM - 1 cm	-13.83	-29.42	1	0.234	0.074
Sta 1, suboxic, POM - 3 cm	0.51	-1.34			
Sta 1, oxic, POM - 1 cm	-2.03	-10.09			
Sta 1, oxic, POM - 3 cm	1.54	0.23			
Sta 7, oxic, DOM - 1 cm	3.05	5.00	-1.3	-1.049	-1.043
Sta 7, oxic, DOM - 3 cm	0.39	0.81			
Sta 7, suboxic, POM - 1 cm	-22.05	-43.75	0.4	-0.829	-1.051
Sta 7, suboxic, POM - 3 cm	0.01	-0.43			
Sta 7, oxic, POM - 1 cm	-4.21	-2.57	0.8	-0.829	-1.051
Sta 7, oxic, POM - 3 cm	0.66	0.40			

**Table S1.**  $\Delta\delta^{13}$ C values of analysed incubations of BHT, BHT' and archaeal lipids (bp = biphytane with 0, 2, or 3 cyclopentane moieties, all from intact polar lipids, and single analysis due to low amounts). The values shown were derived by subtracting the  $\delta^{13}$ C values from background cores from values of incubation cores with <sup>13</sup>C labelled DOM and POM.

			δ <sup>13</sup> C [‰ V-PDB]				
Ту	ре	Sample	bp-0	bp-2	bp-3	тос	
Natural abundance	tural	P900 - 1 cm	-22.6	-21.6	-21.7	-21.5	
	undance	P1800 - 1 cm	-20.4	-19.9	-19.7	-20.3	
		P900, suboxic, DOM - 1 cm	-20.7	-21.8	-21.6	-	
Incubatior		P900, suboxic, POM - 1 cm	-21.6	-21.3	-21.6	-	
	cubations	P1800, oxic, DOM - 1 cm	-21.7	-20.9	-20.7	-	
		P1800, suboxic, POM - 1 cm	-20.0	-20.7	-20.8	-	
		P1800, oxic, POM - 1 cm	-19.6	-20.7	-20.8	-	

**Table S2.**  $\delta^{13}$ C values of analysed samples. bp = biphytane with 0, 2, or 3 cyclopentane moieties, all from intact polar lipids, and single analysis due to low amounts. The values shown are natural abundance from background cores, and from incubations with <sup>13</sup>C labelled DOM and POM, under oxic and suboxic conditions.

#### Text S1. Analysis of archaeal lipids

The values presented in Table S1 were determined as follows:

The freeze-dried subsamples of the background and incubated cores were ground and extracted by a modified Bligh-Dyer extraction method (Lengger et al., 2012b). Briefly, they were extracted ultrasonically three times in a mixture of methanol/dichloromethane (DCM)/phosphate buffer (2:1:0.8, v:v:v) and centrifuged, and the solvent phases were combined. The solvent ratio was then adjusted to 1:1:0.9, v:v:v to separate the DCM phase. Liquid extraction was repeated two more times, the DCM fractions were combined, the solvent was evaporated and the larger particles were filtered out over glass wool. An aliquot was separated into CL and IPL-GDGTs by silica column separation with hexane/ethyl acetate (1:2, v:v) for the CL-fraction and MeOH to elute the IPLfraction.

The IPL fraction was then subjected to ether cleavage in order to release biphytanyl chains from GDGTs (Fig. 1c), following procedures described by Schouten et al. (1998a). To this end, the IPL fraction was refluxed in 57% HI for 1 h to cleave the ether bonds and produce alkyliodides and subsequently extracted 3 times with hexane. The hexane phase was washed with 5% NaS<sub>2</sub>O<sub>7</sub> and twice with water. The alkyliodides were purified over Al<sub>2</sub>O<sub>3</sub> with hexane/DCM 9:1, reduced with LiAlH<sub>4</sub> in 1,4-dioxane for 1 h under reflux, the remaining LiAlH<sub>4</sub> was reacted with ethyl acetate, bidistilled H<sub>2</sub>O was added and the biphytanes were extracted with DCM from the dioxane/H<sub>2</sub>O mixture. Additional purification was achieved by elution over an Al<sub>2</sub>O<sub>3</sub> column using hexane.

GC-MS was used to identify the biphytanes formed upon ether cleavage of GDGTs and PLFAs using a TRACE GC with a DSQ mass spectrometer. The gas chromatograph was equipped with a fused silica capillary column (25 m, 0.32 mm internal diameter) coated with CP Sil-5 (film thickness 0.12  $\mu$ m). The carrier gas was helium. The compound specific carbon isotopic composition of the biphytanes was measured with an Agilent 6800 GC, using the same GC column conditions, coupled to a ThermoFisher Delta V isotope ratio monitoring mass spectrometer. The isotopic values were calculated by integrating the 44, 45 and 46 ion currents of the peaks and that of CO<sub>2</sub>-spikes produced by admitting CO<sub>2</sub> with a known <sup>13</sup>C-content into the mass spectrometer at regular intervals. The performance of the instrument was checked by daily injections of a standard mixture of a C<sub>20</sub> and a C<sub>24</sub> perdeuterated n-alkane of known isotopic composition. Material from the Arabian Sea, outside the OMZ and incubated suboxically with POM yielded insufficient material for  $\delta^{13}$ C determination. The stable carbon isotope compositions are reported in the delta notation against the V-PDB standard.