# Methane transport and sources in an Arctic deep-water cold seep offshore NW Svalbard (Vestnesa Ridge, 79°N)

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   Plouzané, France.
- 14 Abstract.

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15 We investigate the uppermost 60 cm of sediment in active pockmarks of a deep-water methane seep site from

- 16 Vestnesa Ridge offshore NW Svalbard. Using video guided core sampling with a remotely operated vehicle we
- 17 collected push cores directly from bacterial mats within two active pockmarks, Lunde and Lomvi. Pore water
- 18 analyses show very shallow sulphate methane transition zones and transport-reaction modelling suggests a
- 19 considerable amount of dissolved methane passing through the sediment water interface due to upwards
- 20 advection of an aqueous fluid not previously reported from Vestnesa Ridge. In addition, we show that the
- amount of methane that bypasses the benthic methane filter greatly increases with higher aqueous fluid advection
- 22 rate. Recent changes in methane flux are evident from lipid biomarker, seep carbonate, and  $\delta^{13}$ C-organic carbon
- 23 profiles in both pockmarks. Hydrocarbons at this cold seep site are supplied both by deep thermogenic sources
- 24 from below the gas hydrate stability zone but also to a significant degree by microbial methanogenesis which
- 25 dominates the signature in our shallow sediment cores with  $\delta^{13}$ C-CH<sub>4</sub> values as low as -77‰.
- 26 Keywords: methane, aqueous fluid advection, pore water modeling, microbial methanogenesis, Arctic cold seep

#### 27 1 Introduction

- 28 The Arctic is vulnerable to ongoing climate change and potential release of currently trapped geological methane
- in gas hydrates, which might eventually affect the future atmospheric methane concentration (IPCC, 2019).
- 30 However, current knowledge states that seafloor methane release has no significant influence on the atmospheric
- 31 methane concentrations (e.g. Ferré et al., 2020; Myhre et al., 2016) and that anthropogenic climate change is not
- 32 yet enhancing gas-hydrate destabilization (Hong et al., 2017; Wallmann et al., 2018). Studies have focused
- 33 mostly on shallow water seep sites (<400 m) including western Svalbard (Prins Karls Forland) (Jansson et al.,
- 34 2019; Myhre et al., 2016; Silyakova et al., 2020), the East Siberian Arctic shelf (Puglini et al., 2019; Shakhova et

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35 al., 2010) and the Barents Sea Shelf (e.g. Storfjordrenna, Andreassen et al., 2017; Hong et al., 2017). In contrast, 36 seep dynamics from deep-water gas reservoirs are largely unconstrained, except for the Vestnesa Ridge gas 37 hydrate system offshore NW Svalbard. Here, numerous studies have investigated the controlling factors for 38 active methane release from seafloor pockmarks (Knies et al., 2018; Panieri et al., 2017; Plaza-Faverola et al., 39 2015; Schneider et al., 2018; Vogt et al., 1994; Westbrook et al., 2009). Changes in the stress regime due to 40 glacial forebuldge propagation influencing fault activation and permeability (Himmler et al., 2019; Plaza-41 Faverola and Keiding, 2019) have been suggested as factors influencing methane seepage in the past. These 42 constraints on past methane seepage at Vestnesa Ridge are derived from (1) studying paleo-pockmarks in the 43 seismic record (Plaza-Faverola et al., 2015), (2) dating of seep carbonates (Himmler et al., 2019), (3) analysing 44 overgrowth of secondary carbonates on benthic and planktonic foraminifera (Panieri et al., 2016; Schneider et al., 2018) and (4) dating bivalve shell beds with <sup>13</sup>C-depleted signatures which feed on chemosynthetic bacteria 45 (Ambrose et al., 2015). Buried pockmarks and gas chimneys mapped at several stratigraphic intervals in seismic 46 47 profiles suggest episodic seepage activity starting at approximately 2.7 Ma (Plaza-Faverola et al., 2015). Seep 48 carbonate dating suggests that episodic methane seepage in pockmarks on the eastern segment of Vestnesa Ridge 49 has been ongoing since at least the penultimate glacial maximum ~160 ka ago (Himmler et al., 2019). The 50 controlling factor of the present-day seepage pattern, however, is thought to be the orientation of faults in 51 relation to the stress field caused by the nearby ultra-slow mid-ocean ridge spreading (Plaza-Faverola and 52 Keiding, 2019).

53 Numerous studies have investigated gas composition of dissolved methane in the sediment pore water and also 54 in gas hydrate samples (Graves et al., 2017; Panieri et al., 2017; Plaza-Faverola et al., 2017; Smith et al., 2014). 55 However, a limited dataset has hindered reliable conclusions on gas sources in different sediment depths and 56 transformation processes until recently, also because dissolved methane sampled by the headspace method can 57 have undergone intensive biogeochemical modifications already. In a study by Pape et al. (2020) the 58 investigation of gravity cores and MeBo cores from the active Lunde pockmark on Vestnesa Ridge down to a 59 depth of 62 mbsf sheds light on the origin of light hydrocarbons in gas hydrate and sediments on Vestnesa 60 Ridge. The authors find that deep thermogenic hydrocarbons migrate upwards from a depth of more than 800 m 61 and dominate the gas in gas hydrates. Evidence for microbial methanogenesis is found mostly in samples from 62 depths shallower than 50 mbsf suggesting that microbial methane contribution is more important at Vestnesa 63 Ridge than previously thought. However, at higher methane flux sites the signature of microbial methane is 64 mostly overprinted by deeper thermogenic methane.

While studies show the importance of gas migration, the significance of aqueous fluid advection and its impact on local carbon cycling is less emphasized. Hong et al. (2018) showed that aqueous fluid advection could dominate the fluid system once the free gas supply has terminated. Previous studies have investigated how the amount of methane escaping the seafloor varies as a function on fluid flow velocity or different permeabilities (Luff et al., 2004; Stranne et al., 2019; Wallmann et al., 2006), but this has not been investigated at Vestnsea Ridge yet.

71 Here we present pore fluid and sediment geochemistry data from shallow (down to 60 cmbsf) push cores sampled from two active pockmarks using a video guided remotely operated vehicle (ROV). We sampled cores 72 73 directly from locations that exhibit gas discharge features allowing to study the fluid flow regime (i.e. diffusive 74 or advective), and its influence on carbon cycling and the efficiency of the benthic methane filter, using reaction-75 transport modelling. Furthermore, we investigate the gas composition (thermogenic vs microbial) of the 76 dissolved gas in the sediment pore water, and also the relationship of carbon isotopes between the different 77 reservoirs of the cold seep system: methane, dissolved inorganic carbon, sedimentary organic matter and seep 78 carbonate nodules, and assess biogeochemical processes occurring in seep-dominated marine sediments.

#### 79 2 Geological Setting

80 Vestnesa Ridge is a submarine, elongated, c. 100 km long sediment drift located offshore NW Svalbard with an 81 extensive gas hydrate and free gas system (Fig. 1). Vestnesa Ridge lies to the east of the slow-spreading Molloy 82 Ridge, and to the north of the Molloy fracture zone (Hustoft et al., 2009). The sediment succession is several 83 kilometres thick and subdivided into three seismo-stratigraphic sequences (Eiken and Hinz, 1993). The topmost 84 sediments on the ridge crest are late Weichselian and Holocene deposits composed of silty turbidites and muddy-85 silty contourites (Howe et al., 2008). Gas hydrates and seabed fluid flow systems are widespread along continental margins (Judd and Hovland, 2007) but Vestnesa Ridge is a unique setting in that it is situated close to 86 87 a mid-ocean spreading centre and thus is underlain by relatively young and hot oceanic crust with a heat flux of 88 more than 115 mW/m<sup>2</sup> (e.g. Engen et al., 2008; Hustoft et al., 2009). Vestnesa Ridge bends from an E-W 89 direction to a SE-NW direction, and an elongated pockmark field with more than 100 individual pockmarks up 90 to 600 m in diameter are aligned along the crest (Hustoft et al., 2009; Vogt et al., 1994) (Fig. 1). On the western 91 segment pockmarks are scattered on a larger area of the ridge, whereas on the eastern segment they align almost 92 perfectly on the narrow ridge crest. Only the eastern segment has shown periodic active gas discharge in recent 93 years (Bünz et al., 2012).

The bottom-simulating-reflector (BSR) is located at approximately 200 mbsf (Hustoft et al., 2009). Below the BSR, a free gas layer of approximately 30-100 m thickness was inferred (Bünz et al., 2012; Goswami et al., 2015; Hustoft et al., 2009). The topography of Vestnesa Ridge and the prominent BSR anticline below the ridge crest control the gas accumulation. Seismic chimneys, which pierce the complete gas hydrate stability zone (GHSZ), are believed to be the gas migration pathways that supply gas to the seafloor and created the pockmarks (Bünz et al., 2012).



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- 101 Figure 1:A) Overview map of Svalbard and the Fram Strait. B) Bathymetry of the NW Svalbard margin modified from
- 102 (Schneider et al., 2018). Bathymetry data from (Jakobsson et al., 2012). The red square on Vestnesa Ridge is the
- approximate location of the pockmarks Lunde and Lomvi. C) ROV-obtained high resolution bathymetry of the pockmarks
   Lunde and Lomvi with push core locations indicated as red dots and black dots (PC008 from Yao et al., 2019). The dotted
- 105 circle indicates the area of MeBo Seep site SW investigated by (Pape et al., 2020).

## 106 3 Study Area

#### 107 **3.1 Lunde and Lomvi pockmarks**

108 Two of the most active pockmarks in the eastern segment of Vestnesa Ridge are the Lomvi and Lunde

109 pockmarks (Panieri et al., 2017). Seafloor images of these two pockmarks reveal authigenic seep carbonates and

- 110 extensive chemosynthetic communities (Himmler et al., 2019; Panieri et al., 2017). The two pockmarks show
- 111 complex internal bathymetry comprising depressions and ridges (Fig. 1C). 3D seismic data revealed a vertical
- acoustic chimney structure beneath Lund and Lomvi which extends from the base of the hydrate stability zone
- up to the seafloor (Bünz et al., 2012). Reflection anomalies beneath the two pockmarks indicate upward
- advection of free gas and some enhanced reflections could be the result of gas hydrate and/or seep carbonate
- horizons in the subseafloor (Himmler et al., 2019; Panieri et al., 2017).

#### 116 4 Sample collection

117 On research vessel "G.O. Sars" expedition P1606 to the Vestnesa Ridge in July 2016 we collected five push

118 cores (PC007, PC009, PC017, PC020 and PC021) using ROV Ægir 6000. Using video guidance, we chose sites

- 119 close to bacterial mats and exposed seep carbonates in both pockmarks to collect the push cores (Fig. 2, Table 1).
- 120 One push core (PC020) was collected in a location without any signs of seepage (Table 1). Several push cores
- 121 were retrieved during a single ROV dive and stored in a basket at the seafloor until the end of the ROV dive.

122 Immediately upon recovery of the basket the push cores were sampled for pore water using the rhizon technique

123 and for gas using the headspace technique. Headspace gas was sampled using cut off 5 ml syringes and

extracting exactly 3 ml of sediment through pre-drilled holes in the liner. The sediment was transferred to serum

- 125 glass vials containing a 2.5% NaOH solution. They were then crimp sealed and stored at 4°C. Pore water was
- 126 extracted using 5 cm rhizon samplers (Rizosphere, Netherlands) (Seeberg-Elverfeldt et al., 2005), which were
- 127 inserted through pre-drilled holes at intervals ranging from 2 cm to 6 cm. Pore water was collected in 10 ml acid-
- 128 washed syringes. Pore water splits were prepared for alkalinity and dissolved Fe<sup>2+</sup> analyses onboard, and for
- anion, cation (1vol% HNO<sub>3</sub>) and  $\delta^{13}$ C-DIC analyses and nutrient (frozen) analyses in onshore laboratories. After
- 130 offshore sampling, the push cores were frozen onboard and kept at -18°C until further processing.

Table 1:Push core locations, water depth, core length and basic seafloor description of the sampling site. We differentiate
 between black sediment patches (often the precursor of bacterial mats) and white sulfur-oxidising bacterial mats (e.g. Treude
 et al., 2009)

Core	Pockmark	Latitude	Longitude	Water Depth	Core length	Seafloor Observations
		(°N)	(°E)	(m)	(cm)	
PC020	Lunde	79.0075	6.8990	1206.7	72	Muddy brown sediment
(non-seep)						

PC017	Lunde	79.0078	6.8994	1205.0	51	Muddy sediment with black sediment patches
PC021	Lunde	79.0075	6.8989	1206.5	39	Muddy sediment with black sediment patches and some white sulfur-oxidising bacterial mats. Core taken between black sediment patches.
PC007	Lomvi	79.0023	6.9225	1204.0	38	Core from medium-sized white bacterial mat
PC009	Lomvi	79.0025	6.9220	1205.1	27	Area of white bacterial mat (with smaller black sediment patches) among dark brown silty clay, surrounded by seep carbonate crusts; core taken inside white bacterial mat.

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# 135 **5 Methods**

# 136 5.1 Porewater analyses

<u>Total alkalinity</u> (TA) was determined with a pH-controlled titration to a pH just under 4 with an HCl titrant (12M
 Sigma-Aldrich TraceSELECT HCl diluted to 0.012M). The pH electrode was calibrated against pH 4, 7 and 10

139 Metrohm Instrument buffers. We titrated 0.1 to 1 ml of sample aliquot depending on the amount of pore water

140 recovered. About 10 ml of 0.7M KCl was added to each sample to ensure that the pH electrode was fully

submerged. Titrant acid was added while constantly stirring in an open 50 ml beaker. The amount of acid and pH

142 was manually recorded during each addition. TA was calculated from the Gran function plots, which were made

143 by plotting Gran functions against the titrant volume. Gran function is defined as:

144  $(V_0+V_t)$ \*10-pH, where  $V_0$  is the initial volume of sample and  $V_t$  is the volume of titrant added. The TA was then

estimated from the slope and intercept of the regression line from the Gran function plot. Six to ten points were

used for regression. All titrations were done less than six hours after the syringe disconnected from the rhizons.

147 <u>Dissolved iron</u> (Fe<sup>2+</sup>) was determined photometrically using a Shimadzu UVmini-1240 UV-Vis

148 Spectrophotometer at a wavelength of 565 nm. Addition of a commercial ferrospectral solution (prepared

149 onboard with O<sub>2</sub>-free MiliQ water) to the pore water samples causes the formation of a violet colour complex

150 (Collins et al., 1959). The UV absorption at 565 nm is proportional to the concentration of dissolved iron in the

sample (Beer-Lambert law). Calibration was performed using five standards of known Fe<sup>2+</sup> concentration up to

152 1.5 ppm. If samples had higher  $Fe^{2+}$  concentration than 1.5 ppm they were diluted with O<sub>2</sub>-free ascorbic acid

- 153 solution. For sulphate (SO<sub>4</sub><sup>2-</sup>) analyses, we used a Dionex ICS 1100 Ion Chromatograph with an AS-DV
- autosampler and an IonPac AS23 column (eluent: 4.5 mM Na2CO3/0.8 mM NaHCO3, flow: 1ml/min) at the
- 155 Geological Survey of Norway. The relative standard deviations from repeated measurements of different

- 156 laboratory standards are better than 0.5% for concentrations above 0.1 mM. <u>Cation</u> concentrations were
- determined by inductively coupled plasma atomic emission spectroscopy (ICP- AES) at the Geological Survey
- 158 of Norway using a Perkin Elmer Optima 4300 Dual View. Samples were diluted 1:40 before analysis.

159 Methane ( $CH_4$ ) concentration in the sediment was analysed by equilibrium partitioning and measured at the Swiss Federal Institute of Aquatic Science and Technology (EAWAG) using an Agilent Gas Chromatograph 160 161 G1530N with a flame ionization detector. The relative standard deviation of the CH<sub>4</sub> measurements based on 162 repeated measurements of a calibration standard was  $\pm 2.9\%$ . The CH<sub>4</sub> concentration was calculated by adding 163 the headspace CH<sub>4</sub> concentration to the CH<sub>4</sub> concentration in the fluid/sediment which was calculated using the 164 Bunsen coefficients from Yamamoto et al. (1976) taking into account temperature and salinity. Assuming a constant porosity of 0.7 we calculated the CH<sub>4</sub> concentration in the pore water. The porosity value was estimated 165 166 from porosity data from core JR211-26 from Vestnesa Ridge (Hong et al., 2016). Due to potential degassing 167 during core recovery, the measured methane concentrations are always regarded as a minimum value. The 168 isotopic composition of DIC in pore water ( $\delta^{13}$ C-DIC) was determined on the CO<sub>2</sub> liberated from the water after 169 acidification with phosphoric acid. Measurements were performed at EAWAG using an IRMS (Isotope Ratio 170 Mass Spectrometer) from Micromass (Isoprime) equipped with a Gilson 222XL Liquid Handler and a Multiflow 171 from Isoprime. The standard deviation of the  $\delta^{13}$ C-DIC measurements from repeated measurements of a standard was  $\pm 0.12\%$  (1 $\sigma$ , n=52). The stable carbon isotope values for DIC are reported in the conventional  $\delta$  notation in 172 permil (‰) relative to V-PDB (Vienna Pee Dee Belemnite). Concentrations of phosphate and ammonium were 173 174 determined colorimetrically with a Technicon AutoAnalyzer II<sup>TM</sup> component at Oregon State University. Details 175 of the analyses have been documented in the United States Environmental Protection Agency (US EPA) Criteria 176 "EPA 600/4-79-020 Methods for Chemical Analysis of Water and Wastes" and can be found online (EPA, 1983). The stable carbon isotopes of methane ( $\delta^{13}C-CH_4$ ) and ethane ( $\delta^{13}C-C_2H_6$ ) and hydrogen isotopes of 177 178 methane ( $\delta^2$ H-CH<sub>4</sub>) were analysed at Hydroisotop GmbH, Germany, using a GC-MS-IRMS system (Thermo 179 Fisher Scientific GmbH) (for details see Sauer et al. (2015)). The isotopic composition is reported in ‰ (δ-180 values) against the international standards VPDB for carbon and Vienna Standard Mean Ocean Water 181 (VSMOW) for hydrogen.

- 182 **5.2 Sediment geochemistry**
- 183 Core images, X-Ray fluorescence (XRF) and magnetic susceptibility (MS) logging

184 The sediment cores were analysed at the NGU core logging facilities in Trondheim, Norway. In the laboratory 185 the frozen cores were split lengthwise using a bandsaw and a diamond equipped blade. After thawing, the 186 surfaces were cleaned to remove any smeared sediment and to prepare for imaging. High-resolution images were 187 taken using a GeoScan IV colour line-scan camera with an AF Nikkor 50 mm f/1.8D lens mounted to a Standard 188 MSCL (MSCL-S) core logger (GeoTek Ltd., UK). After imaging the core surface was covered with a 4 µm 189 Ultralene film (Spex Sample Prep, USA) to avoid cross-contamination and XRF and MS measurements were 190 taken incrementally at 0.5~cm resolution along the longest axis in the centre of the core surfaces. XRF down 191 core logging was carried out with a Delta Innov-X portable XRF sensor (Olympus NDT Inc., USA) attached to 192 the MSCL-S core logger. The pXRF sensor is equipped with a 4W Rh Tube anode and Si drift detector. Prior to 193 core measurements the pXRF sensor was standardized and a customised standard sample was stationary 194 measured for sensor-control purposes. Successive measurements with voltage settings between 50 and 15 keV 195 voltage and 20 sec exposure time provided spectra covering chemical elements from Mg to Pb. The spectra were 196 converted to element concentration in ppm using the Compton normalization approach described elsewhere (e.g. 197 Kenna et al., 2011). From the element spectrum measured the ratio of calcium (Ca) to titanium (Ti) 198 concentrations were used for this study. MS was measured using a MS2E sensor and a MS3 (both Bartington 199 Instruments Limited, UK) attached to the MSCL-S core logger with 10 sec exposure time at the same sample 200 depths as in XRF measurements. Results are reported in 10<sup>-5</sup>SI units.

201 Sediment samples were taken after core scanning at a resolution of ~2.5 cm for the analyses of carbon and

202 nitrogen content and  $\delta^{13}$ C-TOC. Carbonate nodules were found exclusively in PC009 and sampled for

203 determination of mineral composition by XRD and  $\delta^{13}$ C analyses.

204 <u>Carbon and nitrogen content (LECO)</u>

205 All sediment samples were freeze-dried, ground and homogenized at the Geological Survey of Norway. Total

206 carbon (TC) and total organic carbon (TOC) content of the sediment samples were analysed using a LECO SC-

207 632. For TC determination, subsamples of 300-400 mg were combusted at 1350°C while measuring the

208 production of CO<sub>2</sub>. For TOC analyses, subsamples of 400-500 mg were pre-treated with 10 vol% HCl to remove

- 209 inorganic carbon and then rinsed with MilliQ water. Subsequently they were analysed in the same way as
- 210 described for TC. Total inorganic carbon (TIC) was then calculated as TIC = TC TOC. All results are given in
- 211 weight percent (wt%) and the standard deviation for the TC and TOC measurements was  $\pm$  0.06 wt%. The lower
- 212 limit of detection was 0.06 wt% and 0.1 wt% for TC and TOC, respectively. Total nitrogen (TN) analyses were
- 213 performed with a LECO FP 628 Nitrogen Determinator. Subsamples of 200 mg, wrapped in tin foil, were

combusted to NOx, which was transformed to  $N_2$  and detected with a thermal conductivity cell. The standard deviation of the measurement was  $\pm 0.01$  wt% and the lower limit of detection was 0.1 wt%. The ratio of organic carbon over nitrogen content (hereafter C<sub>org</sub>/N<sub>tot</sub> ratio) was calculated by dividing the wt% values and the multiplying by 1.167 to obtain the atomic ratio.

218 The carbon isotopic composition of sedimentary organic matter ( $\delta^{13}C - TOC$ ) was analysed by EA-IRMS

219 (Elemental Analyzer Isotope Ratio Mass Spectrometry) at Iso Analytical Ltd, UK. Subsampled were

220 decarbonated with 10vol% HCl prior to analyses with a RoboPrep-CN elemental analyzer coupled to a Europa

221 Scientific 20-20 IRMS. Reference materials used for quality control were: reference standard IA-R001 (wheat

flour,  $\delta^{13}$ C: -26.43‰ V-PDB), reference standards IA-R005 (beet sugar,  $\delta^{13}$ C: -26.03‰ V-PDB) and reference

standard IA-R006 (cane sugar,  $\delta^{13}$ C: -11.64‰ V-PDB). Furthermore, every fifth sample was analysed in

duplicate.

## 225 Mineralogy and $\delta^{13}$ C analyses of lithified carbonate nodules

226 The quantitative mineralogical compositions of weakly lithified nodules (in PC009) were determined by X-ray 227 diffraction (XRD) on homogenized bulk-rock powders. Dried nodules were pulverized using agate planet milling 228 for ~1 minute. Powders were analysed on a Bruker D8 Advance X-ray diffractometer using Cu-Kα radiation (40 kV/40 mA) at 3 to 75° 20 range (0.02° 20 steps; time/step = 1 second). Minerals were identified using the 229 230 Diffrac.EVA 3.1 software and quantified using Rietveld modelling and the TOPAS 5.0 software (2 to 3 wt.% 231 uncertainty). Subsamples of the XRD analysed powders were used for stable carbon analyses. The powders were 232 reacted with anhydrous phosphoric acid in a GasBench II preparation line at University of Tartu, Estonia. The released CO<sub>2</sub> gas was analysed with a Delta V Advantage isotope ratio mass spectrometer (reproducibility for  $\delta^{13}$ C 233 234 was estimated  $\pm 0.2\%$ ).

#### 235 Lipid biomarkers

Sediment samples for lipid biomarker analyses (only on PC021) were sampled with an alcohol-cleaned spatula and wrapped in aluminium foil and stored at -20°C. The total lipid extraction (TLE) was performed on ~20 g of sediment with ultrasonification in four solvents with decreasing polarity. The solvents used for extraction were methanol (MeOH) to dichloromethane (DCM) 2:1, MeOH to DCM 1:2 and two last steps using only DCM. The total lipid extract was saponified with NaOH. Fatty acids were methylated to fatty acid methyl esters (FAMEs) from the resulting solution after the neutral fraction was extracted. The neutral fraction was further separated into hydrocarbons, ketones, and alcohols. The alcohol fraction was derivatized to form trimethylsilyl (TMS) adducts 243 for analysis. Individual lipid compounds were analysed using a gas chromatograph (GC) (Thermo Scientific 244 TRACE™ Ultra), equipped with a capillary column (Rxi-5ms, 50 m, 0.2mmID, 0.33 µm df), using helium as a carrier gas at a constant flow rate of 1 mL per min. The initial oven temperature was set to be 50°C, held for 2 min 245 246 and then increased to 140°C at a rate of 10°C per min and held for 1 min. Finally, the temperature was increased 247 to 300 °C at 4 °C per min. The final hold time was 160 min to analyse lipids with higher boiling points in the 248 hydrocarbon and alcohol fractions. Concentrations were determined by flame-ionization detection (FID) against 249 internal standards. Unknown compounds were identified with a quadrupole mass spectrometry (QMS) unit 250 (Thermo Scientific DSQ II) linked to a gas chromatograph. Similarly, compound-specific stable carbon isotope ratios were determined using a magnetic sector isotope ratio mass spectrometry unit (Thermo Scientific Delta V 251 252 Advantage) coupled to a gas chromatography setup with the same specifications as the above. The  $\delta^{13}$ C values are 253 reported against VPDB with an analytical error of  $\pm 1$ %.

#### 254 **5.3** Pore water modelling

255	The trai	nsport-reaction model used in this work was developed following similar consideration as in Hong et al.				
256	(2014).	(2014). We considered seven dissolved species including the isotopes of DIC ( <sup>12</sup> C-DIC and <sup>13</sup> C-DIC) and CH <sub>4</sub>				
257	( <sup>12</sup> C-CH	$H_4$ and ${}^{13}$ C-CH <sub>4</sub> ) as well as sulphate, calcium, and ammonium. In addition, we included organic matter				
258	(POC)	and authigenic calcium carbonate as well as their isotopic signatures in the model ( <sup>12</sup> C-POC, <sup>13</sup> C-POC,				
259	<sup>12</sup> C-car	bonate, and <sup>13</sup> C-carbonate) to account for the observed changes in isotopic signatures. Diffusion of the				
260	solutes	was accounted for using Fick's law with diffusion coefficients corrected for temperature and tortuosity				
261	(Boudre	eau, 1997). There are two types of aqueous fluid advection in the model. One is the burial of pore fluid as				
262	a result	of sedimentation. The pore fluid burial velocity is calculated based on the reduction of porosity				
263	(Boudre	eau, 1997). Fluid advection also occurs due to changes in pressure with a velocity constrained by the				
264	shape o	f the measured pore fluid profiles. Burial of sediment particles was also included which was also				
265	calculat	ted from porosity profiles (Pape et al., 2020). We considered six groups of reactions in this model:				
266	1.	Net anaerobic oxidation of methane ( <b>AOM</b> ) coupled to sulphate reduction ( <b>SR</b> )				
267		$SO_4^{2-} + CH_4 \rightarrow HCO_3^- + HS^- + H_2O_1$				
268						
269	2.	Back flux reaction of AOM (Yoshinaga et al., 2014) (AOM-CR)				
270	3.	CO <sub>2</sub> reduction (methanogenesis) (CR)				
271	4.	Particulate organic carbon degradation through sulphate reduction (POC-SR) and fermentation (POC-				
272		ME) (ME=methanogenesis)				
273		POC-SR: TOC+ 0.5SO <sub>4</sub> $\rightarrow$ DIC + N/C NH <sub>4</sub>				
274		POC-ME: TOC $\rightarrow$ DIC + N/C NH <sub>4</sub>				
275	5.	Authigenic carbonate formation (carbonate precipitation CP)				
276		DIC + Ca $\leftrightarrow$ Carbonate				

Incorporation of DIC and CH<sub>4</sub> into biomass of anaerobic methanotrophic archaea, ANME, (DIC assimilation: **DIC-ASS**, CH<sub>4</sub> assimilation: **CH<sub>4</sub>-ASS**)

There are isotopic fractionations associated with some of the reactions which were accounted for by slightly differentiating the kinetic constants of the <sup>12</sup>C- and <sup>13</sup>C-reactions (Rees, 1973). Detailed formulation of these reactions as well as the corresponding isotopic fractionation can be found in the supplementary information. We assumed seawater composition for the initial and top boundary conditions. We adopted either a no flux boundary condition (i.e., no exchange of material across the deepest cell in the model frame) or a fixed composition for the lower boundary condition. The choice of different types of lower boundary conditions depends on the conditions of the site and will be discussed later.

#### 286 6 Results

## 287 6.1 Seafloor observations

288 The high-resolution bathymetric maps (Fig. 1C) reveal complex internal structures of the Lunde and Lomvi

289 pockmarks with ca. 20 % of the area within both pockmarks containing bacterial mats and seep carbonates.

290 Cores PC007 and PC009 are from the southwestern sector of Lomvi within a depression just next to a ridge

structure. Both cores were retrieved from white sulfur oxidising bacterial mats (Fig. 2). PC017 and PC021 from

292 Lunde were also retrieved from seafloor showing white bacterial mat traces but mostly black sediment patches

293 (Fig. 2). The seafloor around the location of PC020 showed no signs of bacterial mats or other indications of gas-

charged sediments.



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Figure 2:Seafloor images taken by the ROV during push core recovery. Note the white sulphur oxidizing bacterial mats and
 black sediment patches in all photos except for the non-seep site PC020.

## 298 6.2 Pore water geochemistry

299 Phosphate and ammonium concentrations were low throughout all investigated cores (below 11  $\mu$ M PO<sub>4</sub><sup>3-</sup> and below 30  $\mu$ M NH<sub>4</sub><sup>+</sup>) (Fig. 3, Table S3). The strongest rise in PO<sub>4</sub><sup>3-</sup> concentration was observed in the non-seep 300 core PC020 with an increase from 1  $\mu$ M up to 9  $\mu$ M within the upper 16 cm. Fe<sup>2+</sup> and Mn<sup>2+</sup> release to the pore 301 302 water in the upper sediment could only be observed in cores PC021 and PC020 from the Lunde pockmark (Fig 303 3). Core PC021 shows a very narrow zone of dissolved  $Mn^{2+}$  in the upper 4 cm of the sediment. The non-seep core PC020 showed the most pronounced increases in dissolved Mn<sup>2+</sup> and Fe<sup>2+</sup>. Dissolved Fe<sup>2+</sup> was above 1µM 304 in the upper 26 cm of the core with a peak of 21  $\mu$ M at 10 cmbsf. Dissolved Mn<sup>2+</sup> is present in the pore water 305 throughout core PC020 but shows a peak of 14 µM at 8cmbsf. In PC017 dissolved Mn<sup>2+</sup> is present in the pore 306 307 water below 15 cm.



309Figure 3:Pore water concentration profiles of PC007, PC009, PC017, PC020 and PC021. Sulphate and  $\delta^{13}C$ -DIC profiles310adopted from (Dessandier et al., 2020). The orange line represents the depth at which sulphate drops below detection limit.311In PC020 sulphate concentration is high throughout the core. In PC021 we assume sulphate is depleted just below our

312 *deepest sample, thus the orange line is dashed.* 

Strong decreases in  $SO_4^{2-}$  concentrations were observed in four out of the five studied push cores: PC007 and 313 314 PC009 from Lomvi pockmark, and PC017 and PC021 from Lunde pockmark. Our non-seep core PC020 showed 315 no decrease in  $SO_4^{2-}$  concentration down to a depth of at least 60 cmbsf indicating little sulphate reduction at this 316 site (Fig. 3). Complex sulphate profiles in some of our cores hinder determining the exact depth of the sulphate-317 methane-transition-zone (SMT (e.g. Boetius et al., 2000; Iversen and Jørgensen, 1985)). Thus, we indicate the 318 depth of sulphate depletion in Fig. 3. PC007 showed a linear decrease in sulphate concentration from 17 mM to 319 below detection limit within the upper 9 cmbsf. Sulphate concentration was 3.6 mM at 2 cmbsf in core PC009, 320 followed by a slow decrease until reaching the detection limit at 14 cmbsf. In PC017  $SO_4^{2-}$  concentration decreased from 17 mM at 2 cmbsf to below 1 mM at 12 cmbsf (Fig. 3). In PC021, SO<sub>4</sub><sup>2-</sup> concentrations 321 decreased from 27 mM in the bottom water to 2 mM at 14 cmbsf with another SO<sub>4</sub><sup>2-</sup> peak at 22 cmbsf. 322 Headspace methane measurements confirm the presence of methane in the shallow sediments with up to 11.9 323 324 mM CH<sub>4</sub> in PC009, up to 4 mM in PC017, up to 3 mM in PC021 and below 0.009 mM throughout non-seep core 325 PC020. PC009 showed highest TA values up to 57.7 mM at 16 cm, followed by PC007 (TA up to 50.1 mM) and PC017 (TA up to 46 mM). At non-seep site PC020 TA increased to only 4.4. mM at 60 cm, indicating very 326 327 limited production of TA (Fig. 3); seawater TA is approximately 2 mM. In PC017  $\delta^{13}$ C-DIC values became progressively lower downcore and reached values as low as -40‰. Below 10 cm depth there was very little 328 variation in the  $\delta^{13}$ C-DIC profile of PC017. Also, the profiles of PC007, PC009 showed little variation with 329 depth and overall low values: PC007 (-35.4‰) and PC009 (-28.7‰). PC021 showed a decreasing trend of  $\delta^{13}$ C-330 DIC in the upper 10 cm and values down to -37.6% below that. In non-seep core PC020 the  $\delta^{13}$ C -DIC only 331 decreased down to -10% (Fig. 3). In our non-seep core PC020 dissolved Ca<sup>2+</sup>, Mg<sup>2+</sup> and Sr<sup>2+</sup> remained close to 332 333 seawater concentration throughout the core (average Ca<sup>2+</sup>: 10mM, average Mg<sup>2+</sup>: 49.2 mM, average Sr<sup>2+</sup>: 87.1 µM). In contrast, in cores PC007, PC009, PC017 and PC021 Ca2+ concentrations decreased to between 4 and 5 334 mM, i.e. about half the seawater concentration.  $Sr^{2+}$  concentrations decreased to between 28 and 44  $\mu$ M, i.e. to 335 half or even a third of seawater concentration (Fig. 3). Mg<sup>2+</sup> concentrations did not show a coherent pattern. 336

## 337 6.3 Gas isotopes

We measured methane headspace concentration and methane carbon isotopes ( $\delta^{13}$ C-CH<sub>4</sub> in ‰ VPDB) on altogether 29 samples from four different push cores, PC009, PC017, PC021 and PC020 (Fig. 4, Table S4). The results showed large variability in  $\delta^{13}$ C-CH<sub>4</sub> from -77.4‰ (PC017) to -51.4‰ (PC020).  $\delta^{13}$ C-CH<sub>4</sub> values in PC009 varied between -69‰ and -65.8‰, and in PC021 the two lowest samples were below -72‰, whereas all samples above 18 cmbsf where above -64‰ (Fig. 4A). We obtained methane hydrogen isotopic composition  $(\delta^2$ H-CH<sub>4</sub> in ‰ VSMOW) on three push cores (PC009, PC017 and PC021). The methane concentrations in PC020 were too low for hydrogen isotope determination. The  $\delta^2$ H-CH<sub>4</sub> values were very similar in all three cores and varied between -189‰ and -163‰ SMOW (Fig. 4B).

346



348 Figure 4: A)Methane concentration and  $\delta^{13}$ C-CH<sub>4</sub> profiles from the push cores of Lunde (PC017, PC020, PC021) and Lomvi 349 (PC009) pockmark and B) plot of  $\delta^{13}$ C-CH<sub>4</sub> versus  $\delta^{2}$ H-CH<sub>4</sub> indicating the different methane sources (after Whiticar, 1999).

Gas data for comparison is from the following publications: (Graves et al., 2017; Panieri et al., 2017; Pape et al., 2020;

351 Plaza-Faverola et al., 2017; Sauer et al., 2016). The blue arrow indicates the direction of isotopic fractionation during

352 oxidation.

353 We also determined ethane carbon isotopes ( $\delta^{13}$ C-C<sub>2</sub>H<sub>6</sub>) on three push cores (Fig. 5). The values are similar in all

three push cores ranging between -28.7‰ and -25.2‰.



Figure 5: An ethane vs. methane  $\delta^{13}C$  plot including data from PC009, PC017 and PC021. For comparison we plotted data from Vestnesa Ridge (Plaza-Faverola et al., 2017), from a northern Norwegian fjord and ODP sites offshore SW Africa with a microbial signature (Nakagawa et al., 2003; Sauer et al., 2016), and from known thermogenic gas in the Caspian Basin (Katz et al., 2002).

## 360 6.4 Sediment characteristics

- 361 6.4.1 Organic carbon, inorganic carbon, and total nitrogen content
- 362 We determined total carbon, total organic carbon and total nitrogen on 92 sediment samples from five push cores
- 363 (Table S5). The average TOC content was 1.8 wt% in PC007 and PC017, 1.5 wt% on average in PC009 and
- PC021, and 1.3 wt% in non-seep core PC020 (Fig. 6). TN content varied between 0.13 wt% and 0.29 wt%
- 365 (average 0.21 wt%). The atomic Corg/Ntot ratios varied between 6.7 and 11 with an average of 8.7 and showed
- 366 no specific downcore trends in any of the push cores (Fig. 6).

367



368

Figure 6: Sediment characteristics and geochemical profiles of push cores from Lunde pockmark (PC020, PC017, PC021)
 and Lomvi pockmark (PC007, PC009). From left to right: high resolution core photos, magnetic susceptibility (MS),total
 organic carbon content (TOC), organic carbon/total nitrogen ratio (Corg/Ntot), stable carbon isotope ratio of total organic

372 carbon ( $\delta^{13}C$  -TOC), CaCO<sub>3</sub> content and the Ca/Ti ratio from XRF scans (green line). The orange lines denote the depth of

- 373 sulphate depletion inferred from the pore water profiles. The blue fields indicate intervals with high carbonate content 374 inferred from the Ca/Ti profiles and the occurrence of seep carbonate nodules. The grey vertical bar in the  $\delta^{13}$ C-TOC plots
- 375 marks the "background" value inferred from non-seep core PC020.
- Total inorganic carbon average values varied between 0.5 wt% (PC017) and 2.3 wt% (PC009). CaCO<sub>3</sub> content
- 377 calculated from the inorganic carbon content varied between 3.8 wt% (PC017) and 19.5 wt% (PC009) (Table 2).
- $\delta^{13}$ C-TOC values showed larger differences between cores and specific downcore trends (Fig. 6). The non-seep
- 379 core PC020 showed the least variation in  $\delta^{13}$ C-TOC values, with a range from -24.3‰ to -23.1‰ (average: -
- 23.7%). All other cores showed strong decreases in  $\delta^{13}$ C-TOC downcore with values lower than -30%, with
- 381 lowest values at the base of the cores (PC007: -30.7‰, PC009: -34.5‰, PC017: -30.2‰, PC021: -30.7‰).

Table 2: Average values of total organic carbon (TOC),  $\delta^{13}C$ -TOC, total nitrogen (TN), total inorganic carbon (TIC), and calcium carbonate content calculated from inorganic carbon (CaCO<sub>3</sub>).

			TOC (wt%)	$\delta^{13}$ C-TOC	TN (wt%)	TIC (wt%)	$CaCO_3$ (wt%)
Lunde	PC020	non-seep	1.3	-23.7	0.17	0.8	6.9
Lunde	PC017	seep	1.8	-25.3	0.23	0.5	3.8
Lunde	PC021	seep	1.5	-25.5	0.20	0.8	7.0
Lomvi	PC007	seep	1.8	-27.8	0.24	0.7	5.8
Lomvi	PC009	seep	1.5	-30.3	0.23	2.3	19.5

## 385 6.4.2 Sediment core images, carbonate content and magnetic susceptibility

386 The core images of the five push cores (Fig. 6) show a range in colours and lightness. PC009 and PC021 are

387 much lighter than PC007 and PC017. This difference corresponds well with their different CaCO<sub>3</sub> content (Fig.

6). PC020 shows some distinct reddish-brown to yellowish-brown coloration of the sediments in the top

- 389 centimetres probably indicating the presence of iron oxides (Schwertmann and Fitzpatrick, 1992). The inferred
- 390 presence of iron oxides corresponds well with the peak in magnetic susceptibility in the upper 15 cm of PC020, a

391 general measure for the concentration of magnetic minerals in marine sediments (Dekkers, 1978). All other cores

392 show very little variation in magnetic susceptibility and overall low values (below  $15 \times 10^{-5}$  SI).

# 393 6.4.3 Diagenetic carbonate nodules

384

Figure 6 shows the CaCO<sub>3</sub> wt% (calculated from total inorganic carbon) and the Ca/Ti ratio (from XRF scans,

- green line) for all cores. The two analyses correspond very well in each core, showing the same downcore trends.
- 396 Only in PC021 there is a peak in Ca/Ti at around 35 cmbsf which is not resolved in the discrete CaCO<sub>3</sub>. We
- found indications for carbonate nodules in core PC021 at around 35 cmbsf (peak in Ca/Ti, white coloration of
- the core) and in core PC009 below 10 cmbsf (increasing abundance with depth). In PC009 the proportion of
- 399 calculated CaCO<sub>3</sub> content reaches up to around 40 wt% at the base of the core. Only in this core we found

400 carbonate nodules large and abundant enough to be extracted by sieving for XRD and carbon isotope analyses 401 (Table 3, Fig. 7). The  $\delta^{13}$ C-carbonate values range between -26.6‰ and -29.4‰ and correspond well with the 402 present-day pore water  $\delta^{13}$ C-DIC values at the base of the core (Fig. 7). The carbonate mineralogy is mainly Mg-403 calcite and aragonite, with only minor proportions of calcite (Table 3, Fig. 7). The main lithogenic components 404 were quartz, plagioclase, illite and muscovite (Table 3).



405 Table 3: Results of quantitative XRD analyses and  $\delta^{3}C$ -carbonate analyses of the authigenic seep carbonates from PC009.

408 Figure 7:left:  $\delta^{13}C$  values of pore water DIC (green dots) and of carbonate nodules (black crosses) of PC009 including a 409 colour photo of the core, right: mineralogy from quantitative XRD analysis of the carbonate nodules extracted from PC009.

410 The occurrence of the carbonate nodules in PC009 corresponds very well with the low  $\delta^{13}C$  -TOC values in this

411 core. Also, in Core PC021 a peak in the Ca/Ti record corresponds to the depth of low  $\delta^{13}$ C -TOC values (Fig. 6).

# 412 6.4.4 Lipid biomarkers in PC021

413 PC021 was sampled for the quantification and compound-specific  $\delta^{13}$ C analysis of archaeal lipid biomarkers,

- 414 namely archaeol and sn2-hydroxyarchaeol, which are isoprenoid glycerol diethers. Their compound-specific
- 415  $\delta^{13}$ C values have been widely used as a tool to identify the presence of AOM communities (e.g. Niemann and
- 416 Elvert, 2008, and references therein). The archaeol content showed peaks at around 8 cm and 36-38 cm sediment
- 417 depth with  $0.33 \,\mu$ g/g and  $0.63 \,\mu$ g/g, respectively. These peaks are less pronounced but also visible in the sn2-
- 418 hydroxyarchaeol content (Table 4). The  $\delta^{13}$ C -of archaeol and sn2-hydroxyarchaeol show low values (down to -

- 419 97‰ and -110‰, respectively) in the areas with elevated lipid biomarker content, confirming their link with the
- 420 presence of ANME communities.

421	Table 4: Content of the lipid biomarkers archaeol and sn2-hydroxyarchaeol and their compound-specific $\delta^{l3}C$ values
422	measured on samples from PC021.

Sediment		sn2-	δ <sup>13</sup> C-	$\delta^{13}$ C-sn2-
depth	archaeol	hydroxyarchaeol	archaeol (‰	hydroxyarchaeol
(cm)	$(\mu g/g)$	$(\mu g/g)$	VPDB)	(% VPDB)
2.7	0.09	0.04		
5.3	0.07	0.09	-67.4	-100.4
8.5	0.33	0.13	-78.7	
11.5	0.05	0.06	-90.5	-90.8
14.3	0.06	0.07	-60.0	-59.3
18.5	0.08	0.03	-63.6	
21.0	0.18	0.20	-82.0	-78.3
24.6	0.12	0.06	-68.9	
26.9	0.28	0.14	-66.8	-82.7
29.3	0.16	0.01	-82.9	
31.6	0.13	0.00	-87.6	-71.5
33.8	0.20	0.00	-71.6	-67.7
36.1	0.63	0.22	-90.6	-110.1
38.2	0.41	0.09	-97.3	-99.5

# 423 7 Discussion

#### 424 7.1 Early diagenetic sequence in shallow sediments and fluid flow indications

425 Pore water sulphate concentrations are controlled by two main processes, 1) the degradation of organic matter via SO<sub>4</sub><sup>2-</sup> reduction (POC-SR) and 2) sulphate-dependent anaerobic oxidation of methane (AOM)(e.g. Reeburgh, 426 427 2003). All cores from the Lunde and Lomvi pockmark showed low concentrations of ammonium and phosphate, 428 which suggests low organic matter degradation rates (Froelich et al., 1979). Similar concentrations were reported from cold seeps south of Svalbard and the northern Norwegian margin (Hong et al., 2020). Peaks in Fe<sup>2+</sup> and 429 430 Mn<sup>2+</sup> (as in PC020 and PC021) are commonly also associated with organic matter degradation utilizing Mn and Fe (oxyhydr)oxides as electron acceptors and releasing dissolved  $Mn^{2+}$  and Fe<sup>2+</sup> to the pore water (Froelich et al., 431 432 1979). Supported by the low ammonium and phosphate concentrations we conclude that the sulphate profiles are 433 controlled mainly by ascending methane through sulphate-dependent-AOM. Anaerobic oxidation of methane leads to the production of bicarbonate documented by the strong increase in the alkalinity in our seep cores. 434 435 Carbonate alkalinity can induce the formation of authigenic seep carbonates, as observed in PC009, including calcite, aragonite, high- and low-magnesium calcite (Aloisi et al., 2002; Bayon et al., 2009; Bian et al., 2013; 436 437 Gieskes et al., 2005; Sauer et al., 2017; Teichert et al., 2003).

- Another indicator of AOM observed in all seep cores is the decrease in  $\delta^{13}$ C-DIC to low values which cannot be explained solely by the degradation of organic matter (except in non-seep PC020). It requires the transformation of isotopically light methane to dissolved inorganic carbon to explain  $\delta^{13}$ C-DIC values as low as -40‰ (PC017). Below the SMT, an increase in  $\delta^{13}$ C-DIC has often been observed in other seep settings (e.g. Haese et al., 2003; Hong et al., 2014; Pape et al., 2020; Paull et al., 2000; Sauer et al., 2015), which is commonly attributed to methanogenesis by CO<sub>2</sub> reduction, preferentially consuming <sup>13</sup>C-DIC approaches or maintains low values even
- 445 well below the SMT.



446

447Figure 8: Scatter plot comparing pore water  $\delta^{13}$ C-DIC to total alkalinity (TA) values in different cores. Vestnesa Ridge data:448Pape et al. (2020)(MeBo seep site SW c. 15 m and reference site c. 62 m), and Yao et al. (2019) (PC008, push core c. 35 cm),449Blake Ridge ODP Site 994: Paull et al. (2000) (ODP Hole 994, c. 700 m). The orange star marks seawater values. The450arrows indicate different downcore trends. The total alkalinity values around the SMT at Seep Site SW, Reference Site, and451Blake Ridge range between 15-30 mM.

- 452 To emphasize this unusual downcore  $\delta^{13}$ C-DIC trend in our pore water profiles, we compare the patterns of
- 453  $\delta^{13}$ C-DIC plotted against TA with published data from the Vestnesa MeBo Seep Site SW and reference site
- 454 (Pape et al., 2020) and Blake Ridge (Paull et al., 2000). In Fig. 8 most of the cores show a trend departing from
- 455 near-seawater values (orange star) and subsequently show the expected downcore addition of <sup>13</sup>C-depleted DIC
- 456 from AOM (PC021, Blake Ridge, Vestnesa Seep- and Reference Site) or by organic matter degradation (PC020).

This results in an increase in alkalinity and at the same time lowering of  $\delta^{13}$ C-DIC values (arrow starting at the 457 458 orange star). Below the SMT in the cores from Vestnesa reference and Seep Site SW, and Blake Ridge, a progressive removal of <sup>13</sup>C-depleted DIC can be observed, indicated by the arrow pointing towards high  $\delta^{13}$ C-459 460 DIC values (Fig. 8), explained by CO<sub>2</sub> reduction. In contrast, the push cores from our study and PC008 from Yao 461 et al. (2019) show a different downcore trend, indicated by the arrow pointing to high TA values. Below the SMT, total alkalinity still increases and  $\delta^{13}$ C-DIC values are continuing to decrease slightly or remain close to 462 463 values found at the SMT (Fig. 3 and 8). Assuming TA is only produced from AOM the maximum concentration is expected around the SMT, but some of our cores show higher TA below the current SMT (PC007, Fig. 3). In 464 order to investigate the processes creating the observed  $\delta^{13}$ C profiles in our cores, we applied a transport-reaction 465 model on both the MeBo Seep site SW (Pape et al., 2020) and our core PC017. We first applied the model to the 466 MeBo Seep site SW with a common set of reactions, such as AOM, carbonate precipitation and CO<sub>2</sub> reduction, 467 which have been shown to occur based on the box modelling performed by Pape et al. (2020). Our model was 468 able to reproduce the measured pore water profiles observed at MeBo Seep Site SW (Fig. 9) with the increase in 469 470 DIC concentration in the uppermost meter that can be attributed to sulphate dependent AOM which also 471 stimulates carbonate precipitation as reflected by the drawdown in calcium concentration (see also the rates of AOM and carbonate precipitation in Fig. 9). The downcore decrease in  $\delta^{13}$ C-DIC in the upper meter results from 472 473 <sup>13</sup>C-depleted DIC production during AOM while the associated decrease in  $\delta^{13}$ C-CH<sub>4</sub> within the same interval 474 reflects the back-reaction of AOM in the model which produces  ${}^{13}C$ -depleted CH<sub>4</sub> (Yoshinaga et al., 2014). 475 Below the SMT the progressive increase in  $\delta^{13}$ C-DIC can be modelled by assigning an in-situ methanogenesis 476 (POC-ME in Fig. 9) and a fixed fluid composition for the lower boundary condition at 5 mbsf ([DIC]=18 mM 477 and  $\delta^{13}$ C-DIC=0‰). Such a lower boundary composition reflects both the addition of DIC through organic carbon fermentation and methanogenesis via  $CO_2$  reduction. In terms of  $\delta^{13}C$ -CH<sub>4</sub> below the SMT, little 478 479 downcore variation is predicted by the model except for the depths around the SMT. Such results indicate that 480 the isotopic signatures of methane largely reflect the source composition of methane from depth at this site, also 481 observed by Pape et al. (2020). We show that the data can be satisfactorily fitted with no additional fluid flow 482 included, only advection due to sediment burial (Fig. 9). This agrees with the findings from Pape et al. (2020) 483 that these pore water profiles can be accounted for with a box model which assumes no advection.



484

485Figure 9: Modelling results for MeBo Seep Site SW (Pape et al., 2020) (pore water advection is only due to sediment burial486and compaction, no additional fluid flow). The total fluid flow in the lower left graph is  $3.02x10^{-4}$  m/yr. The black dots487represent measured pore water composition (Pape et al., 2020), the solid lines represent the best fit from our model. The488upper right graph includes measured data for  $\delta^{13}$ C-DIC (filled dots) and  $\delta^{13}$ C-CH4 (circles). The lower right graph includes489the depth profiles of the rates of organoclastic sulphate reduction (R-POC-SR) and organic matter degradation by490fermentation (R-POC-ME). Rates of AOM, POC-SR and POC-ME are all expressed in mol DIC/m<sup>3</sup>/yr to make them easier to491compare.

492 We then applied our model to PC017, the longest push core besides the non-seep core, showing the least

493 complex pore water profiles. In core PC017 the depth of sulphate depletion is around 13 cmbsf whereas it is at

494 ~110 cm at MeBo Seep site SW. We find that the only way to fit the measured pore water profiles was to 1)

495 reduce the number of reactions included in the model, 2) assign a high fluid advection rate and 3) assign a fixed

496 composition for the lower boundary condition. We tested two other model setups with no external fluid

497 advection (Fig. S1), and with neither CO<sub>2</sub> reduction nor external fluid advection (Fig S2), but found

498 unsatisfactory fits for DIC, Ca<sup>2+</sup> and  $\delta^{13}$ C-DIC profiles in both tests (supplementary information).

In Fig. 10 we show that the pore water profiles, especially the DIC concentration and  $\delta^{13}$ C-DIC pattern can only

500 be explained when including fast fluid advection (0.2 m/yr) of a fluid containing no sulphate, abundant DIC and

501 methane (40 mM and 6 mM, respectively, both with relatively low  $\delta^{13}$ C values), low concentrations of

502 ammonium (20 µM) as well as calcium (5 mM). This composition of the ascending fluid is used as the lower

503 boundary condition. A fraction of the DIC is consumed by CO<sub>2</sub> reduction (see R-CR in Fig. 10), but it does not

- significantly influence the overall downcore  $\delta^{13}$ C-DIC signature, i.e. there is only little increase in  $\delta^{13}$ C-DIC
- with depth. The minimal increase in  $\delta^{13}$ C-DIC with depth in PC017 can be explained by the influence of the
- solution ascending fluid which dominates the  $\delta^{13}$ C-DIC signature. The amount of methane produced by CO<sub>2</sub> reduction

507 (with low  $\delta^{13}$ C-CH<sub>4</sub> values) is however enough to cause a local minimum in  $\delta^{13}$ C-CH<sub>4</sub> values in the PC017

profile at around 15 cm where the highest rate of  $CO_2$  reduction is estimated (Fig. 10). This result means that the

509 addition of methane through  $CO_2$  reduction is significant enough, compared to the concentration of methane

510 transported in the ascending fluid, to influence the overall  $\delta^{13}$ C-CH<sub>4</sub> signature.



511

Figure 10: Modelling results for PC017 with a total fluid advection rate of 0.2 m/yr. The black dots and circles represent
measured pore water data, the solid lines represent the best fit of our model. The lower right graph includes the depth
profiles of the rates of CO<sub>2</sub> reduction (R-CR) organoclastic sulphate reduction (R-POC-SR) and organic matter degradation
by fermentation (R-POC-ME), and organic matter formation by the AOM microbial consortium (R-POC formation). Rates of
AOM, POC-SR, POC-ME and R-CR are all expressed in mol DIC/m<sup>3</sup>/yr to make them easier to compare.

517

518 The phase of the fluid and mode of transport are critical in terms of the effectiveness of the biological benthic 519 methane filter, i.e. how the microbial community can utilize the methane. Dissolved methane can be transported via diffusion and/or by advection of the pore water itself. Even though diffusion of methane is ubiquitous and 520 521 occurs wherever there is a concentration gradient, there is usually very limited amounts of methane reaching the 522 sediment-water interface in diffusion-dominated environments due to the efficient AOM (e.g. Egger et al., 2018). 523 Advection of aqueous fluid with high dissolved methane concentration, however, can transport considerable 524 amounts of methane towards the sediment-water interface (e.g. Luff et al., 2004; Stranne et al., 2019). Gas 525 advection can also cause large amounts of methane to escape from the sediment through fractures or channels (MacDonald et al., 2002). Previous studies have shown that Lunde and Lomvi pockmark are characterized by 526 diffusion, focussed fluid flow and methane transported in the gas phase (Hong et al., 2016; Panieri et al., 2017; 527

528 Pape et al., 2020; Yao et al., 2019). With our model we show that at site PC017 it requires a fast-moving 529 aqueous fluid with dissolved methane to explain the observed pore water profiles. Given the constant  $\delta^{13}$ C-DIC profiles we suggest that cores PC007, PC009, and PC008 (Yao et al., 2019) were dominated by aqueous fluid 530 531 advection as well. High methane concentration and low  $\delta^{13}$ C-DIC downcore variation in PC009 indicates especially efficient aqueous fluid advection. Worth noticing, our model predicts that only 6 mM of dissolved 532 533 methane in the ascending pore water is enough to induce the observed change in fluid composition, a methane 534 concentration that is far lower than what is required to form gas hydrate given the 1200m water depth. Pape et al. 535 (2020) documented the presence of gas hydrate as shallow as 45 cmbsf from Vestnesa Ridge, and also at 536 Storfjordrenna south of Svalbard, gas hydrate was observed at 35 cmbsf (Klasek et al., 2020). The shallow SMT 537 at our investigated sites, but the absence of gas hydrate in all of our cores seems to support the relatively lower 538 concentration of dissolved methane derived by our model. 539 In addition, we conducted a sensitivity test by varying the total fluid advection rate to investigate how the fluxes 540 of the different solutes in the pore water, especially methane, vary (Table 5). This is done based on the model 541 results from PC017. At a total advection rate of 0.2 m/yr, which is needed to reproduce our pore water profiles, 542 67% of methane entering our sediment core at the bottom of the core will pass the sediment water interface and 543 escape into the water column. This is in stark contrast to MeBo Seep Site SW where only 1% of methane 544 entering at the bottom of the core will leave the sediment into the water column. Our results are in agreement 545 with previous modelling studies of methane transport across the sediment water interface (Stranne et al., 2019; 546 Wallmann et al., 2006) and again emphasize the critical role of aqueous fluid advection..

547 Table 5: Model results of methane flux at the sediment water interface and at the bottom of the core (PC017) depending on
548 the advection rate. The last line shows the results from MeBo Seep Site SW.

total advection rate	flux at the seafloor	flux at the bottom of the core	
m/yr	CH4/m2/yr	CH₄/m₂/yr	% of CH₄ leaving sediments
0.2	0.57	0.85	67%
0.15	0.34	0.64	53%
0.1	0.17	0.45	37%
0.05	0.06	0.33	18%
0.01	0.02	0.38	6%
0.001	0.02	0.41	4%
0.0001	0.02	0.42	4%
MeBo Seep Site SW	0.00	0.23	1%

550 At certain sites within the pockmarks Lunde and Lomvi methane is bypassing AOM/the benthic methane filter, 551 otherwise there would be no acoustic gas flares detected in the water column (e.g. Panieri et al., 2017). Methane gas is not available for microbial utilization, therefor the microbial filter is always damned to fail when gas 552 553 migrates (e.g. Luff and Wallmann, 2003; Treude et al., 2003). Furthermore, Yao et al. (2019) showed that gas 554 advection along mini-fractures can rapidly increase methane concentrations in shallow sediments and play an 555 important role in modulating methane dynamics in shallow sediments. However, our push cores allow for the 556 first time to show and quantify considerable dissolved methane escape from the sediment at Vestnesa Ridge 557 based on pore water profiles. Two studies that have investigated pore water profiles from Vestnesa Ridge by 558 modelling (Hong et al., 2016; Pape et al., 2020) both suggest that at the investigated sites the AOM benthic 559 methane filter is effectively preventing dissolved methane from entering the water column. In contrast, our push 560 cores show that a significant amount of dissolved methane is passing the sediment-water interface (Table 5). 561 This also highlights the value of core sampling with video guidance, in this case the use of an ROV, which 562 allows to specifically target sites with methane seepage indicators (e.g. bacterial mats).

# 563 7.2 Organic matter origin and past changes in methane flux

564 Both the content and  $\delta^{13}$ C signature of TOC in our push cores are more extreme compared to previously published data from Vestnesa Ridge. Pape et al. (2020) report TOC values ranging from 0.51 to 1.27 wt% at 565 566 MeBo Seep site SW (Lunde pockmark) whereas in our push cores from seep locations we measured TOC as high as 2 wt% (Fig. 6). Furthermore,  $\delta^{13}$ C-TOC values are as low as -34.5‰ in PC009, such low values have not 567 568 been reported from Vestnesa Ridge previously. Similar  $\delta^{13}$ C values of organic matter extracted from a seep site (-33‰) have been reported from the Eel River Basin offshore northern California (Hinrichs et al., 1999). 569 570 Common tools to determine the origin of organic matter in sediments are the Corg/Ntot ratio and  $\delta^{13}$ C-TOC 571 values. However, this approach is limited to regions where simple mixing between marine and terrestrial organic 572 matter is the case. At Vestnesa Ridge the use of Corg/Ntot ratios and  $\delta^{13}$ C-TOC values to determine organic 573 matter sources is complicated by the contribution of seep-endemic methanotrophic organisms. We interpret the negative trends in  $\delta^{13}$ C-TOC in our push cores as a contribution from microbial biomass consisting of 574 575 methanotrophic archaea and sulphate reducing bacteria rather than from terrestrial organic carbon that has an 576 average  $\delta^{13}$ C-value of -27‰ (e.g. Rachold and Hubberten, 1999). This is also supported by the Corg/Ntot values, which do not show excursions to higher values where  $\delta^{13}$ C -TOC values are low (Fig. 6), which would be 577 578 expected from terrestrial organic carbon. We assume our  $\delta^{13}$ C -TOC values to represent mixing between marine

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579 organic matter (i.e. primary production) and methanotrophic/chemosynthetic biomass which seem to have 580 similar Corg/Ntot ratios, but large differences in  $\delta^{13}$ C values. All push cores, except for the non-seep core, showed strong decrease in  $\delta^{13}$ C-TOC with depth, which is interpreted as an increasing proportion of 581 methanotrophic biomass with depth. PC009 with the lowest  $\delta^{13}$ C-TOC values presumably contains the highest 582 proportion of methanotrophic biomass. Also, PC021 shows a pronounced decrease in  $\delta^{13}$ C -TOC from around 14 583 584 cm (-23.4‰) to 38 cm (-30.7‰). When compared with our lipid biomarker data, from 17 cm and downward, an 585 increasing trend in archaeol content and a decreasing trend in  $\delta^{13}$ C-archaeol and  $\delta^{13}$ C-sn2-hydroxyarchaeol can be deduced (Fig.11). This pattern agrees well with the decreasing trend of  $\delta^{13}$ C of the TOC pool and confirms 586 587 that this is caused by the presence of methanotrophic microbial biomass. There is another interval further towards the top of PC021 (c. 8 cmbsf) with elevated archaeol content and relatively lower  $\delta^{13}$ C values of both 588 589 archaeol and sn2-hydroxyarchaeol. This could indicate SMT fluctuations as marked by the grey intervals in Fig. 590 11. Another possibility could be a situation similar to the double SMT observed offshore north-western Svalbard 591 by Treude et al. (2020). The authors suggest non-vertical methane transport causing a second interval with an 592 activity maximum of sulphate dependent AOM.



593

- 594 Figure 11: from left to right: pore water sulphate and methane concentration in PC021, the  $\partial^{3}C$  of total organic carbon
- 595 (TOC) in PC021, and the content and compound specific  $\delta^{13}C$  of the lipid biomarkers archaeol and sn2-hydroxyarchaeol. 596 The grey bands represent intervals with elevated content of archaeol and relatively low values of  $\delta^{13}C$  of both archaeol and 597 sn2-hydroxyarchaeol.
- To determine whether the decrease in  $\delta^{13}$ C-TOC in our seep push cores can be explained by the incorporation of
- <sup>13</sup>C depleted carbon from methane or DIC during AOM we included -organic carbon formation in our transport-
- 600 reaction-model (see reaction rate R-POC formation in Fig 10). Previous studies show that ANME can assimilate
- 601 carbon from both methane and DIC (Kellermann et al., 2012; Wegener et al., 2008), and that usually very low
- amounts of carbon are assimilated compared to amounts of methane oxidised (e.g. around 1 mol%, Wegener et.,

603 al 2008). The result of our modelling exercise for PC017 shows that the highest rate of organic carbon formation 604 occurs at around 12 cm, very close to the SMT, where AOM rates are highest (Fig. 10). However, the lowest 605  $\delta^{13}$ C-TOC values in PC017 are found at the bottom of the core at 38 cmbsf. We consider two different options to 606 explain this discrepancy. In the first option, the SMT with the highest rates of AOM and carbon assimilation by 607 the AOM-performing microbial community might have been situated further down in the sedimentary column previously. The fact that  $\delta^{13}$ C-TOC values in all push cores, except the non-seep core (PC020), are lowest below 608 609 the present-day SMT supports this explanation and indicates a lower methane flux in both Lunde and Lomvi 610 pockmark in the past. For the second option, both the low  $\delta^{13}$ C-TOC values and the seep carbonates in PC009 611 below the present-day SMT could, on the other hand, also be explained by a higher methane flux in the past. 612 When these sediment layers were situated near the sediment surface under higher flux conditions this would 613 have resulted in the accumulation of <sup>13</sup>C-depleted TOC and aragonite- and Mg-calcite-dominated seep 614 carbonates close to the sediment water interface. A subsequent burial to grater depths (where they are found 615 now) and a lowering in the methane flux could explain the absence of seep carbonates in our push cores close to 616 the sediment surface and  $\delta^{13}$ C-TOC values closer to background values at the top of the cores (Fig. 6).

No matter whether an increase or a decrease in methane flux was responsible for the geochemical patterns in our push cores, from our data set we cannot ascertain the reason for a change in fluid flow. However, cold seeps are known to show very dynamic fluid escape patterns. Previous studies suggest that the regional stress regime is the main external factor controlling the long-term pattern of gas seepage at the Vestnesa Ridge (Plaza-Faverola and Keiding, 2019). Shorter-term changes in fluid flow might be due to local gas hydrate destabilization caused by ascending warmer fluids, changing fluid pathways due to local clogging of fractures by gas-hydrates or seep carbonates (e.g. Himmler et al., 2019; Luff et al., 2004), or by the formation of new fractures (Treude et al.,

624 2020; Yao et al., 2019)

#### 625 7.3 Microbial methanogenesis fuelled by thermogenic hydrocarbon-sustained biomass

626 The large variability in  $\delta^{13}$ C-CH<sub>4</sub> values in our cores is probably the result of the mixing of different proportions

627 of thermogenic and microbial methane and the influence of AOM. The effect of methane oxidation was

especially visible in PC020 where methane concentrations in the sediments were low throughout the core (<9

 $\mu$ M.). This methane represents the residual after oxidation and is thus enriched in <sup>13</sup>C compared to methane

- 630 sampled below the SMT, which has not been strongly affected by oxidation yet. PC017 showed the lowest  $\delta^{13}$ C-
- 631 CH<sub>4</sub> values between -77.4‰ and -73.3‰ indicating the highest proportion of microbial methane (Whiticar,

632 1999). Combining carbon and hydrogen isotopic data of methane (Fig. 4B), can help interpret the methane

633 source (Whiticar, 1999). All our samples from cores PC009, PC017 and PC021 plot in the field of

634 hydrogenotrophic methanogenesis (CO<sub>2</sub> reduction), which is the process of methanogenic archaea growing on

carbon dioxide and using  $H_2$  as a source of electrons to reduce  $CO_2$  (Richards et al., 2016). Four samples from

636 PC021 plot close to the mixing field of microbial and thermogenic methane but these samples might be affected

by methane oxidation, which increases the  $\delta^{13}$ C-CH<sub>4</sub> value (as indicated by the arrow in Fig. 4B). Although

638 methane isotopes seem to indicate a microbial origin,  $\delta^{13}$ C-C<sub>2</sub>H<sub>6</sub> values are well within the thermogenic range

(Fig. 5). The plot of  $\delta^{13}$ C-C<sub>2</sub>H<sub>6</sub> vs.  $\delta^{13}$ C-CH<sub>4</sub> values shows that gas samples of known microbial origin show

 $\delta^{13}$ C-C<sub>2</sub>H<sub>6</sub> values mostly below -60‰ (Nakagawa et al., 2003), more than 30‰ lower than in the Vestnesa push core samples.

642 There is evidence supporting the hypothesis that the gas hydrate system and large-scale fluid migration at 643 Vestnesa Ridge is fuelled by deep thermogenic hydrocarbons. The gas analyses of gas hydrate samples show 644 exclusively thermogenic signatures (Graves et al., 2017; Panieri et al., 2017; Pape et al., 2020; Plaza-Faverola et 645 al., 2017; Smith et al., 2014)(Fig. 4). Seismic chimneys, consisting of vertical zones of disrupted reflections, 646 connect to active pockmarks and are rooted in the free gas zone below the BSR (Hustoft et al., 2009). The 647 location of the chimneys and pockmarks on the crest correlates with the location of the BSR anticline and is also 648 related to the occurrence of fault planes which can facilitate fluid migration (Hustoft et al., 2009; Plaza-Faverola 649 et al., 2015). Hydrocarbon migration modelling suggests large scale fluid migration started around 6 million 650 years ago and reached the seafloor around 2 million years ago (Knies et al., 2018). Miocene source rock 651 observed in ODP Hole 909 and inferred below Hole 909 (Knies and Mann, 2002; Stein et al., 1995) have been 652 suggested to fuel the gas hydrate system at Vestnesa Ridge since then. This is also discussed in detail by Pape et 653 al. (2020) who show that thermogenic hydrocarbons dominate the gas signature at the seep sites including the 654 gas hydrate samples and point towards a common deep-hydrocarbon source for both Lunde and Lomvi 655 pockmark. On the other hand, shallow gas samples from our push cores in Lunde and Lomvi pockmark sampled 656 at seep locations show a strong signal of microbial methane which had not been observed from Vestnesa Ridge yet. We suggest two possible ways to explain this: (1) microbial methanogenesis in shallow sediment as a result 657 658 of organic matter degradation fuelled by methane-seep related biomass or (2) secondary methanogenesis as a result of thermogenic hydrocarbon biodegradation (e.g. Milkov, 2011; Mishra et al., 2017; Stagars et al., 2017). 659 660 Some admixture of thermogenic gas in our samples is evident from the high ethane  $\delta^{13}$ C values (Fig. 5). But due to a lack of  $\delta^{13}$ C measurements of higher hydrocarbons it is at this stage not possible to conclude on possible 661 662 biodegradation of the higher hydrocarbons. In both scenarios of microbial methane formation, the thermogenic

663 hydrocarbon seepage is a prerequisite to produce microbial methane in the shallow subsurface sediments, either through providing higher hydrocarbons for biodegradation, or providing seep habitats that induce accumulation 664 665 of organic matter. The pockmarks are a result of large-scale hydrocarbon migration and produce methane seeps 666 that are a biological oasis on the high Arctic seafloor. For example, seep carbonates can enhance the occurrence 667 of fauna, because they provide sites for aggregation. Also fractures or pockets in the sediment often provide 668 space and nutrients for microbial habitats that create a favourable condition for higher biomass accumulation 669 such as biofilms (Briggs et al., 2011; Gründger et al., 2019). A study by Åström et al. (2018) shows that the 670 infaunal biomass is 5 times higher at the Vestnesa pockmarks than at a high Arctic deep-sea control site. 671 Furthermore, our sedimentary organic matter analyses suggest considerable methanotrophic biomass in the 672 sediment. Another prerequisite for microbial methanogenesis is low sulphate concentrations. Sulphate is 673 removed rapidly in shallow sediments due to sulphate dependent AOM as shown by our pore water profiles. The 674 high supply of organic matter and the low concentrations of pore water sulphate can thus facilitate microbial 675 methanogenesis (e.g. Paulo et al., 2015, and references therein), which shows that this can only occur as a result 676 of the thermogenic hydrocarbon migration. This is supported by the fact that in our non-seep core PC020, which 677 showed no seep indications, there is no shallow microbial methane detected, although TOC content is relatively 678 high.

#### 679 8 Conclusions

1. Pore water profiles from seep sites within Lunde and Lomvi pockmark show numerous indications of anaerobic oxidation of methane occurring in shallow subsurface sediments, such as rapid sulphate depletion, strong increase in alkalinity and low  $\delta^{13}$ C-DIC values. The depth profiles of  $\delta^{13}$ C-DIC and alkalinity, combined with transport reaction modelling suggest advective aqueous fluid transport assists the escape of methane from the Lunde pockmark sediments. This results in considerable flux of dissolved methane across the sediment-water interface not previously reported from Vestnesa Ridge and highlights the value of video guided sampling of seep sites which had not been done at Vestnesa Ridge previously.

2. The depth profiles of  $\delta^{13}$ C-TOC and lipid biomarker data indicate a significant amount of AOM-associated microbial biomass in our sediment cores that do not correspond to the present-day sulphate-methane transition. The negative trends in  $\delta^{13}$ C-TOC and the occurrence of seep carbonates below the present-day SMT could be explained either by a previously deeper SMT in the sediment and thus a lower methane flux in the past, or by a previously higher methane flux creating strong <sup>13</sup>C-depletion in the TOC and seep carbonate accumulation in near-surface sediments which were subsequently buried to deeper depths preserving the SMT-signature. 693 3. The large-scale fluid flow and gas-hydrate system at Vestnesa Ridge is fuelled by deep thermogenic

694 hydrocarbons and we observed a clear thermogenic signature in the  $\delta^{13}$ C of ethane gas in our push cores.

695 However, distinct signals of microbial methane in the shallow subseafloor sediments are also evident from our

696 push cores. We link this to microbial methanogenesis which is sustained only by the occurrence of the seeps that

697 provide habitats for the increased production of biomass (the basis for methanogenesis), compared to the

- 698 'background' Arctic seafloor where organic carbon content is lower and no methane is found in the shallow
- 699 subsurface.

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