constance.choquel@gmail.com on behalf of the coauthors

Dear Editor and referees,

Please find the new version of the paper "Denitrification by benthic foraminifera and their contribution to N-loss from a fjord environment". The main corrections performed to the manuscript are highlighted in pink.

Briefly, the corrections requested related to improving the quality of the English manuscript. This manuscript was proofread by 3 English-speaking coauthors.

- The Abstract of the manuscript has been clarified and improved.
- The Introduction and the Material and Methods have been lightened and English improved, while keeping the content remains unchanged with all comments from reviewers taken into account.
- The internet address of SMHI has been included in the bibliographic list.
- Few corrections were made on the Results only to clarify the reading.
- A clarification was made on Figure 5 and in the text. Indeed, the first denitrification zone of the GF17-3 station is 1.2 to 3.5 cm deep.
- Four bibliographic references suggested by the coauthors have been added.

For more details on minor changes please refer to the manuscript with corrections.

Best regards. Constance Choquel

Denitrification by benthic foraminifera and their contribution to N-loss from a fjord environment

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Abstract.

Oxygen and nitrate availabilities impact the marine nitrogen cycle at a range of spatial and temporal scales. Here, we demonstrate the impact of denitrifying foraminifera on the nitrogen cycle at two oxygen and nitrate contrasting stations in a fjord environment (Gullmar Fjord, Sweden). The Denitrification by benthic foraminifera contribution to benthic denitrification was stimated by coupling living foraminifera determined through the combination of specific density countings per microhabitat denitrification rule measurement and sedimentary specific nitrate respiration rates obtained through incubation experiments using N2O microsensors. Benthic nitrate removal was calculated from submillimeter chemical gradients extracted from 2D distribution porewater images of the porewater nitrate concentration. These were acquired combining diffusive equilibrium the DET technique (Diffusive Equilibrium in the Hins-(DET Thin film) with chemical colorimetry and hyperspectral imagery. Oxygenated bottom waters Sediments with high nitrate contentconcentrations in sediment porewatershe porewater and oxygenated overlying water were dominated by the non-indigenous species (NIS) Nonionella sp. TI which T1. Denitrification by this species could denitrify up toaccount for 50-100 % of nitrate porewater. Contributions in the nitrate gradients. In contrast sediments below hypoxic bottom waters where sediment porewaters were nitrate had low inventories of porewater nitrate, and denitrifying foraminifera were scarce and did not contributerare. Their contribution to nitrogen mitigation in oxic coastal ecosystems and should be included in ecological and diagenetic models aiming at understandingto understand biogeochemical cycles coupled to nitrogen.

1 Introduction

Hypoxic water (i.e. [O₂] < 63 μmol L⁻¹ Diaz et al., 2008; Breitburg et al., 2018) occurs frequently in bottom-waters of shallow coastal seas, due to remineralization of organic matter and water stratification. In this study we used the hypoxia threshold of 63 μmol L⁺(e.g. Diaz et al., 2008; Breitburg et al., 2018). Hypoxia may have large ecological effects (Levin et al., 2009; Rabalais et al., 2010; Zhang et al., 2010), such as an increase of fauna mortality (Stachowitsch et al., 1984; Diaz et

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al., 2001). However, certain microorganisms, e.g. bacteria and foraminifera, can perform denitrification by respiring nitrate (Risgaard-Petersen et al., 2006) and thereby survive in depleted oxygen environments.

The effects of decreasing dissolved oxygen availability at spatial and temporal scales will impact biogeochemical cycles such as the nitrogen cycle (Childs et al., 2002; Kemp et al., 2005; Conley et al., 2007; Diaz et al., 2008; Neubacher et al., 2013; Breitburg et al., 2018). The nitrogen cycle in marine sediments is a perpetual balance between nitrogen inputs (e.g. terrestrial runoff, atmospheric precipitations) and outputs (e.g. denitrification from sediment and water column) (Galloway et al., 2004; Sigman et al., 2009). In most semi-enclosed marine environments as the Baltic Sea, the nitrogen loss through benthic denitrification exceeds the inputs of nitrogen through nitrogen fixation. These nitrogen sink regions of the ocean are mostly associated with anoxic regions (Gruber and Sarmiento 1997). This study focuses on how one important compartment of the marine meiofaunal community—the benthic foraminifera—is coupled to the nitrogen cycle during contrasted dissolved [O₂] conditions at two different stations, focusing on the impact of a non-indigenous species (NIS).

The nitrogen cycle occurring in marine sediments is dependent on the bottom-water oxygenation. In<u>At</u> oxic bottom water conditions (Fig. 1a), ammonium (NH₄⁺) produced from remineralization of particulate organic nitrogen (PON) in sediments, diffuses toward the oxic sediment-superficial layer and through the <u>water</u>-sediment<u>-water</u> interface. (SWI). Nitrification ean occur is an aerobic process which converts NH_4^+ to nitrate (NO_3^-) in the oxic sediment and in the oxic water column through the conversion of NH_4^+ to nitrate (NO_3^-) (Rysgaard et al., 1994; Thamdrup and Dalsgaard, 2008). Conversely, Total denitrification occurs in sediment when oxygen is scarce (below 5 µmol L^+ , Devol et al., 2008) and organic earbon and nitrate are available. Denitrification named, the sum of "canonical denitrification" ($NO_3^- \rightarrow NO_2^- \rightarrow NO \rightarrow N_2O$ $\rightarrow N_2O$ and anammox is an anaerobic process whereby nitrate is used as the terminal electron acceptor in the oxidation of organic matter by facultative anaerobic metabolisms when oxygen is exhausted. Denitrification participates in the loss of the fixed Nitrogen to N_2 gas, that converts NO_2^- or NO_3^- to N gasses, such as e.g. N_2 (Brandes et al., 2007 and references within). Another process can contribute to this loss of N_2 gas: Anammox (anaerobic ammonia oxidation) (Engström et al., 2005; Brandma et al., 2011). According to Brandes et al. (2007 and references within) the "total denitrification" can be defined as the sum of the canonical denitrification plus the anammox. Nitrification and denitrification are thus strongly coupled, and

denitrification can be enhanced by adjacent sedimentary nitrification zones or by direct NO₄ diffusion) generating N removal from the overlying water towards the sedimentenvironment. The (Kemp et al., 1990; Cornwell et al., 1999). When bottom water turns hypoxic, the nitrogen cycle occurring in the sediment is strongly affected (Fig. 1 b). Nitrate production is reduced since nitrification cannot process under low oxygen conditions typically occurs in sediment layers where oxygen is scarce (i.e. < 5 µmol L⁻¹, Devol et al., 2008) and (~ 0 µmol L⁻¹; Rysgaard et al., 1994; Mortimer et al., 2004). However, deeper into reduced sediment, nitrification can occur through secondary reactions with NH₄ oxidation by Mn and Fe oxides (Luther et al., 1997; Mortimer et al., 2004). Denitrification is the dominant process of nitrate reduction in coastal marine sediments. (Thamdrup and Dalsgaard, 2008; Herbert, 1999). Denitrification depends on the nitrate transported from the water column and adjacent sedimentary nitrification zones. Nitrification and denitrification are thereby strongly coupled (Kemp et al., 1990; Cornwell et al., 1999). This dependency on nitrification can imply a reduction of denitrification rates as bottom water turns hypoxic, (Fig. 1 b) since nitrification rates are reduced as nitrification cannot proceed under low oxygen concentrations (~0) umol L⁻¹; Rysgaard et al., 1994; Mortimer et al., 2004). The exception however is anoxic nitrification occurring through secondary reactions with NH₄⁺ oxidation by Mn and Fe oxides (Luther et al., 1997; Mortimer et al., 2004). However, dissimilatory nitrate reduction to ammonium (DNRA) can also contribute to nitrate depletion in reduced sedimentIn reduced sediment, dissimilatory nitrate reduction to ammonium (DNRA) can also contribute to nitrate depletion leading to NO₃conversion into NH₄⁺ instead of nitrogen (N₂) (Christensen et al., 2000) and compete denitrification.

Benthic foraminifera were the first marine eukaryotes found to perform <u>complete</u> denitrification (Risgaard-Petersen et al., 2006), but not all foraminifera species can denitrify (Piña-Ochoa et al., 2010). Denitrifying foraminifera species are defined in our study as species able to perform denitrification proved by denitrification rate measurements. <u>TheseThe</u> denitrifying species have a facultative anaerobic metabolism and <u>store</u> nitrate—<u>storing foraminifera in their cells, which</u> can <u>use</u> either environmental oxygen or nitrate to respire (Piña-Ochoa et al., 2010). <u>be used for denitrification.</u> Nonionella cf. stella (Charrieau et al., 2019 and references therein) and Globobulimina turgida were identified as the first denitrifying foraminifera species (Risgaard-Petersen et al., 2006). <u>Currently</u>) but <u>currently</u>, nineteen denitrifying species <u>within 9 genera</u> are known (Glock et al., 2019). <u>Foraminifera denitrification rates show a large</u>Their cell specific rate range from 7 ± 1 pmol N indiv-

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d⁻¹ to 2241 ± 1825 pmol N indiv₇⁻¹ d⁻¹ (Glock et al., 2019), and the contribution of benthic foraminiferal communities to benthic denitrification lies in the range from 1 to 90 % (Kamp et al., 2015, Dale et al., 2016; Xu et al.,)-2017).

Recently, a non-indigenous and suspected invasive *Nonionella stella* morphotype: *Nonionella* sp. T1 was described as invasive in the North Sea region (Deldicq et al., 2019) and also reported infrom the Gullmar Fjord (Sweden) (< 5 %, Polovodova Asteman and Schönfeld, 2015). The genus *Nonionella* is potentially capable to denitrify as demonstrated for *Nonionella* cf. *stella*, by Risgaard-Petersen et al. (2006). However, the NIS *Nonionella* sp. T1 morphotype, differs both morphologically and genetically from *Nonionella stella* specimens sampled inpreviously at other localities, such as the Santa Barbara Basin (California USA) differs morphologically (Charrieau et al., 2018) and genetically (Deldicq et al., 2019) from the specimens sampled in-), the Kattegat and OslofjordOslo Fjord (Norway), respectively.-) (Deldicq et al. (., 2019) describe these specimens as the *Nonionella* sp. T1 morphotype, a non-indigenous and suspected invasive species in the Oslofjord. The genus *Nonionella* is potentially capable to denitrify as demonstrated with *Nonionella* cf.-). As a consequence, the denitrification capacity *stella* by Risgaard Petersen et al. (2006). Denitrification rates of two species from the NIS *Nonionella* sp. T1 is unclear.

In the present study, we investigate if the suspected invasion of the NIS Nonionella sp. T1 has any implication for the nitrogen cycle in sections of the Gullmar Fjord have been measured(Sweden) that is subjected to hypoxic events. Several denitrifying foraminifera species are present in the Gullmar Fjord sediments: Globobulimina turgida (Risgaard-Petersen et al., 2006) and), Globobulimina auriculata (Woehle et al., 2018). Additionally, Stainforthia fusiformis and Bolivina pseudopunctata are two dominant species in the deepest part of the fjord (Gustafsson and Nordberg, 2001; Filipsson and Nordberg, 2004). These species are also potential candidates for The denitrification. Indeed, capacity of the denitrification rates of Stainforthia fusiformislatter two species in the Gullmar Fjord is indicative from Perú were measured by Piña Ochoa et al. (2010) and severaldirect measurement on affiliated species sampled at the coast of Bolivina from PerúPeru, Bay of Biscay (France) and Santa Barbara were measured by Basin (Glock et al. (2019); Piña-Ochoa et al. (2010) and Bernhard et al. (2012), respectively. On the other hand, other typical fjord species such as (2012). Several species, which apparently lack the ability to denitrify, but are able to survive anoxia, are, however, also present in the sediments of the fjord. These include Bulimina marginata, Cassidulina laevigata, Hyalinea balthica are considered as non-denitrifying species by Piña Ochoa et al. (2010) as their intracellular nitrate reserves are almost absent. The anaerobic metabolism of some other species commonly found in

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the fjord such as Leptohalysis scotti, Liebusella goesi, Nonionellina labradorica and Textularia earlandi-is not documented in previous studies. In the context of ecosystem function and service, it is therefore of interest if the NIS Nonionella sp. T1 can denitrify and thereby if its invasion into the Gullmar Fjord maintains (or elevates) the denitrification capacity of the overall foraminifera community and thus the sediment or, alternatively, if the organism share a metabolism similar to the non-denitrifying specimens above, with the possible consequence that the suspected invasion of NIS Nonionella sp. T1 implies reduced contribution of foraminifera based denitrification to the loss of N from the fjord.

A high abundance of denitrifying foraminifera in both oxic and anoxic marine environments play an important role in the nitrogen cycle (Risgaard-Petersen et al., 2006; Piña-Ochoa et al., 2010; Bernhard et al., 2012; Glock et al., 2013; Xu et al., 2017). Previous estimates of foraminifera contributions to denitrification range from 1 to 90 % (Dale et al., 2016; Xu et al., 2017). Estimates of foraminifera contribution to benthic denitrification are limited by the high spatial and temporal variability of sediment geochemistry and distribution of denitrifying foraminifera, which poses particular methodological challenges. Marine sediments often include chemical micro-heterogeneities (Aller et al., 1998; Stockdale et al., 2009), which can be averaged out within the volume of a sediment slice. Moreover, sediment core slicing or centrifugation can induce cell lysis, which can induceded to a bias in porewater nitrate concentrations (Risgaard-Petersen et al., 2006). To characterize theseTo obtain better estimates of the chemical microenvironments at relevant submillimeter/ millimeter scales, new approaches have to be used. Recently, a 2D-DET (two Dimensions Diffusive Equilibrium in Thin-film) technique combining ombined with colorimetry and hyperspectral imagery was developed to obtain the distribution of nitrite and nitrate in sediment porewater at millimeter resolution in two dimensions (Metzger et al., 2016). This method avoids mixing of intracellular nitrate and the nitrate contained in the sediment porewater. We will apply this technique here to get information about the distribution and concentration of nitrate at a scale relevant for modeling denitrification rates.

The present study aims to examine how the NIS Nonionella sp. T1 and the other denitrifying species affect the nitrogen cycle by comparing two stations with contrasting oxygen and nitrate environments subjected to hypoxic events.

Thegeneral objectives of the paperstudy are: (1) to characterize the density of the living benthic foraminifera at two contrasted stations; (2) to measure the denitrification rate contrasting stations in the Gullmar fjord: one with oxic bottom water and one

with hypoxic bottom water. We will in particular focus on the relative abundance of the NIS *Nonionella* sp. T1 (2) to investigate if this NIS *Nonionella* sp. T1 can denitrify and (3) to quantify its eventual contributions to benthic denitrification; (4) to in the sediments. On the basis of the results we will discuss the probable future impact of the NIS *Nonionella* sp. T1 on the foraminifera fauna and the nitrogen cycle in the Gullmar Fjord.

2 Material and Methods

2.1 Site description and sampling conditions

The Gullmar Fjord is 28 km long, 1-2 km wide and located on the Swedish West coast (Fig. 2). The fjord undergoes fluctuations between cold and temperate climates (Svansson, 1975; Nordberg, 1991; Polovodova Asteman and Nordberg, 2013; Polovodova Asteman et al., 2018). The fjord is stratified (Fig. 2 d) in four water masses (Svansson, 1984; Arneborg, 2004). Hypoxia events in the fjord have been linked to the influence of the North Atlantic Oscillation (NAO) (Nordberg et al., 2000; Björk and Nordberg, 2003; Filipsson and Nordberg, 2004). Several monitoring stations are located in the fjord: Släggö (65 m water depth), Björkholmen (70 m water depth) and Alsbäck (117 m water depth), the hydrographic and nutrient data were obtained from the SMHI's publically (Swedish Meteorological and Hydrological Institute)Institute's (SMHI) publicly available data-base SHARK (Svenskt HavsarkivSMHI, 2020). Since 2010, the threshold of hypoxia ([O₂] < 2 mg L⁻¹, i.e. 63 μmol L⁻¹) in Alsbäck station (red squares, Fig. 3) is reached typically in late autumn and winter. Deep-water exchanges usually occur in late water-early spring. However, the duration of hypoxia varies between years and hypoxia events also-occurred in the summer 2014 and 2015, due to lack of deep-water exchange. The frequency of hypoxic events has increased in the fjord (Nordberg et al., 2000; Filipsson and Nordberg, 2004).

Two sampling cruises were conducted in the Gullmar Fjord on board R/V Skagerak and Oscar von Sydow, respectively. The 2017 cruise (GF17) took place between 14th and 15th November 2017 and two stations were sampled (GF17-3 and GF17-1, Fig. 2 c and d) to define the living foraminifera fauna and the sediment geochemistry at two contrasted stations. The 2018 cruise (GF18) took place on the 5th September 2018 with the focus to collect living Nonionella sp. T1 for O₂ respiration and denitrification rates measurements. Only one station (at the same position as GF17-3) was sampled.

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GF17-3 (50 m water depth) is located closest to the mouth of the fjord (58°16′50.94″N/ 11°30′30.96″E) with bottom waters from Skagerrak (blue diamond, Fig. 3) and GF17-1 (117 m_water depth) close to the deepest part of the fjord (58°19′41.40″N/11°33′8.40″E) near Alsbäck monitoring station in the middle of the stagnant basin (red square, Fig. 3). In November 2017, CTD profiles indicated the water mass structures at both stations (Fig. S1). Bottom water at GF17-3 station was oxic with a dissolved oxygen content of 234 μmol L⁻¹. The dissolved oxygen content decreased strongly with depth at the GF17-1 station reaching 9 μmol L⁻¹ at the seafloor, which is below the severe hypoxia threshold.

2.2 Foraminifera sampling and processing

During the 2017 cruise, two sediment cores per station (1A, 1C and 3A, 3C for GF17-1 and GF17-3 stations respectively) were immediately subsampled with a smaller cylindrical core (\emptyset 8.2 cm) and sliced every 2 mm upfrom the sediment surface to 2 cm depth and every 5 mm from 2 cm to 5 cm depth to study living foraminifera distribution. The samples were incubated without light for 10–19 hours in ambient seawater with Cell Tracker Green (CMFDA, 1 mM final concentration) at *in situ* temperaturestemperature (Bernhard et al., 2006) and then fixed with ethanol 96°. Fixed samples were sieved (> 355, 150, 125 and 100 μ m) and the > 100 μ m fraction, the most commonly fraction used for foraminiferal analyses in the Gullmar Fjord (see Charrieau et al., 2018 and references therein) was examined using an epifluorescence microscope equipped for fluorescein detection (i.e., 470 nm excitation; Olympus SZX13). In the present study, the foraminifera distribution will be described highlighting the NIS *Nonionella* sp. T1.

2.3 Geochemical sampling and processing

One core from the shallow GF17-3 station was reserved for O_2 microelectrode profiling. Oxygen concentration was measured in the dark with a Clark electrode (50 μ m tip diameter, Unisense ®, Denmark) within the first 5 mm depth at a 100 μ m vertical resolution. Due to technical problems, no oxygen profiling was done at the GF17-1 station.

One core per station was dedicated for geochemical analyses; they were carefully brought to Lund University (Sweden) and stored at the sampling site in situ temperature (10°C) until further analysis the next day. Overlaying water of the GF17-3 core was gently air bubbled to maintain the oxygenated conditions recorded at this station. Overlaying Hypoxia in the

overlaying water of the GF17-1 core was bubbledmaintained by bubbling with N₂ gas passed through a solution of carbonate/bicarbonate to avoid pH rise due to degassing of CO₂-by N₂-bubbling.

Nitrite/Nitrate were analyzed using the 2D-DET method from Metzger et al. (2016). In brief, for each core, a DET (Diffusive Equilibrium in Thin films) gel probe (16 cm x 6.5 cm and 0.1 cm thickness) was hand-made prepared. The gel probe was inserted into the sediment and left for 5 hours to allow for a diffusive equilibration time-between the gel and porewaters. After equilibration, the gel was removed of the core and laid on a first NO₂⁻ reagent gel. After 15 mminutes at ambient temperature thea pink coloration must appear werewhere nitrite is detected. A reflectance analysis photographimage of the nitrite gels fauna was taken with a hyperspectral camera (HySpex VNIR 1600). The next step was to convert existing nitrate into nitrite with the addition of a reagent gel of vanadium chloride (VCl₃). After 20 minminutes at 50°C, additional pink coloration is interpreted as -porewater nitrate concentration. Followed by the acquisition of another hyperspectral image and convertedthe conversion into false colourscolors through a calibrated scale of concentrations, the final image wasgel images were cropped to avoid border effects. Each pixel (190 μm x 190 μm) was decomposed as a linear combination of the logarithm of the different end-member spectra using ENVI software (unmixing function) (Cesbron et al., 2014; Metzger et al., 2016). Nitrite and nitrate detection limit is 1.7 μmol L⁻¹ (Metzger et al., 2016). Nitrate production/consumption zones for each station were estimated by extracting the average and standard deviation of the 290 vertical 1D profiles ((5.5 cm width x 1 pixel) / 0.019 cm for 1 pixel size) on the 2D gels and modelling using PROFILE software (Berg et al., 1998)) limits are 1.7 μmol L⁻¹ (Metzger et al., 2016).

2.4 Oxygen and nitrate respiration and denitrification rates rate measurements of the NIS Nonionella sp. T1

The two cores sampled <u>induring</u> the 2018 cruise (GF18) at the shallower GF17-3 station were carefully transported <u>and stored</u> at *in situ* temperature (8 °C) and stored for three days at the Department of Geosciences, Aarhus University (Denmark). *Nonionella* sp. T1 specimens were picked <u>underat</u> *in situ* temperature and collected in a Petri dish, containing a thin layer of sediment (32 µm) to check their vitality. Only living, active *Nonionella* sp. T1 specimens were picked <u>using a brush</u> and cleaned several times <u>using a brush</u> with micro-filtered, nitrate-free artificial seawater.

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Oxygen respiration rates were measured, following the method developed by Høgslund et al. (2008) using a Clark type oxygen microsensors (50 μ m tip diameter, Unisense ®, Denmark) (Revsbech, 1989)). The O₂ sensor was calibrated by a two-point calibration using air-saturated water at *in situ* temperature (8 °C) and sodiumin 0.7 M alkaline ascorbate solution (to strip O₂ out of the system) as zero. O₂) and air-saturated sea water. Then, a pool of 5five living *Nonionella* sp. T1 was transferred into a glass microtube (inner diameter 0.5 mm, height 7.5 mm)), that was fixed inside a 20 ml test tube mounted in a glass-cooling bath (8 °C). A motorized micromanipulator was used to measure O₂ concentration profiles along a distance gradient that ranged from 200 μ m of the foraminifera to 1200 μ m using 100 μ m steps. Seven O₂ concentration profiles were generated with one incubation containing the pool of *Nonionella* sp. T1. Negative controls were done by measuring O₂ rates from microtube with empty foraminifera shells and blanks with empty microtube. Oxygen respiration rates were calculated with Fick's first law of diffusion, J = -D * dC/dx, where J is the flux, dC/dx is the concentration gradient obtained by profiles and D is the free diffusion coefficient of oxygen at 8 °C for a salinity of 34 (1.382 x 10.5 cm. Ramsing and Gundersen, 1994). The seven O₂ respiration rates were calculated as the product of the flux by the cross section area of the microtube (0.196 mm²). Then, the average O₂ respiration rate was divided by the number (n=5) of *Nonionella* sp. T1 presented present in the microtubes to obtain the respiration rate per individual.

The same pool of *Nonionella* sp. T1 specimens as for the O₂ respiration ratesmeasurements was used for denitrification rate measurements. Denitrification ratesThese measurements were measuredperformed in the microtubes as it is described in Risgaard PetersenHøgslund et al., (2006). In this method, denitrification is stopped at the N₂O production by acetylene inhibition that can be measured with a (2017). A N₂O microprobe (Andersen et al., 2001) with a 50 µm tip diameter, Unisense @, Denmark). Nitrous oxide was measured as used to measure the end product instead of N₂ (Risgaard Petersen et al., 2006). Nitrous oxide flux was estimated from the chemical gradient profiled from the pool of *Nonionella* sp. T1 inserted in a microchamber. The N₂O production was multiplied by two because two moles of NO₃-are required for the production of one

microchamber. The N_2O production was multiplied by two because two moles of NO_3 ⁻ are required for the production of one mole of N_2O (Risgaard Petersen et al., 2006). The microchamber is porous to gases and is bathed in a sodium ascorbate solution that maintains oxygen N_2O concentration at zero within-profile, that developed in the chamber after acetylene inhibition of the final step in the denitrification process ($N_2O \rightarrow N_2$). Calibration of the microchamber. The microchamber was filled with an

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5 mM of Hepes buffer (to maintain the pH stable). Calibrationsensor was performed using the standard addition method by successive injections of a N₂O saturated solution in order to have 14 μM steps of final concentration. Negative controls were done by checking the absence of O₂ from microchamber with empty foraminifera shells and blanks with empty microchamber. Then, the pool of Nonionella sp. T1., was transferred to the microchamber with a micropipette. The N₂O The cell specific N₂O production rate was calculated from the N₂O flux (estimated from the concentration profiles were repeated seven times on gradient and Fick's first law), the surface area of the microtube (0.25 mm²) and the pool number of Nonionella sp. T1. The source of nitrate during denitrification comes from intracellular nitrate storage of Nonionella sp. T1 (not measured in this study)-the tubes (n=5) as described above. Rates are reported with the unit pmol N indiv⁻¹ d⁻¹.

Since O_2 respiration and denitrification rates are linked to cytoplasmic volumes or biovolumes (BV) (Geslin et al., 2011; Glock et al., 2019), the specimens from the pool of *Nonionella* sp. T1 were measured (width (a) and length (b)). Fig. 4) using a micrometer mounted on a Leica stereomicroscope (MZ 12.5) to estimate the average BV. The volume of the shellseach shell was estimated by using the best resembling geometric shape, a spheroid prolate $(V = \frac{4}{a^3} \pi (\frac{a}{2})^2 (\frac{b}{2}))$. Then, according to Hannah et al., (1994)). 75 % of the measured entire volume of the shell was used corresponding to as the estimated cytoplasmic volume. To compare the size of the Five Nonionella sp. T1 specimens sampled induring the 2017 cruise (GF17, study of the fauna) were also measured to compare their average size with the Nonionella sp. T1 samples insize of the specimens sampled during the 2018 cruise (GF18, denitrification rate measurements), 5 specimens sampled in the 2017 cruise were also measured.)

2.5 Contributions of the NIS Nonionella sp. T1 to diffusive oxygen and nitrate uptakebenthic denitrification.

The following estimated contributions to sediment diffusive oxygen and nitrate uptake were performed mainly on the dominant denitrifying species, *Nonionella* sp. T1.Benthic denitrification was estimated using the 2D nitrate concentrations obtained with the DET technique. An average 1D nitrate profile was obtained by calculating the mean of 290 vertical profiles ((5.5 cm width x 1 pixel) / 0.019 cm for 1-pixel size) extracted from the 2D concentration image. Then, nitrate production and consumption zones were calculated with PROFILE software (Berg et al., 1998). With the assumption that nitrate consumption

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was equivalent to denitrification, the benthic denitrification rate was calculated by integrating nitrate consumption over the depth.

The denitrification activity of the NIS *Nonionella* sp. T1 population was calculated using the specimen abundances in the nitrate consumption zones and their cell specific activity. The size of the *Nonionella* sp. T1 specimens sampled during the two cruises, however, differed markedly (Table 1). We need to correct the denitrification rate of *Nonionella* sp. T1 specimens from the 2017 cruise to take into account the difference of shell size. The measured *Nonionella* sp. T1-cell specific denitrification rate (2018 cruise) was normalized by specimen BV (2017 cruise) using of denitrifying foraminifera is correlated with their size according to the relationshipmodel; $\ln (y) = 0.68 \ln (x) - 5.57$, where y is the denitrification rate (pmol ind dai) and x is the shell BV (μ m) ((Geslin et al., 2011; Glock et al., 2019; Equation S1). The corrected *Nonionella* sp. T1 denitrification rate is multiplied by the *Nonionella* sp. T1 specimens counted found in each denitrifying zones defined by PROFILE modelling. Then, two calculation approaches were discussed to and we therefore used this model to correct the denitrification estimates for size specific variations.

A maximum estimate of the contribution of the NIS Nonionella sp. T1 contributions population to benthic denitrification:

(A) to divide the—was obtained from the ratio of the denitrification activity of Nonionella sp. T1 denitrification rate by the nitrate porewater population and the benthic denitrification rate estimated from PROFILE modelling, then the second calculation (B) to divide the Nonionella sp. T1 denitrification rate by the total denitrification from PROFILE plus the Nonionella sp. T1 denitrification rate. In the first approach (A) we suggest Nonionella sp. T1the porewater nitrate concentration profiles. This presumes that Nonionella sp. T1 exclusively use only the nitrate dissolved in the sediment porewater. In the second approach (B) we suggest that the foraminifera use both intracellular and—as source for denitrification (calculation approach A). A minimum estimate of the contribution of Nonionella sp. T1 population to benthic denitrification was obtained from the ratio between the denitrification activity of Nonionella sp. T1 population and the benthic denitrification rate estimated from porewater nitrate pools for denitrification. concentration profiles plus the denitrification activity of Nonionella sp. T1 population. This presumes that Nonionella sp. T1 exclusively use intracellular nitrate as source for denitrification (calculation approach B).

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3 Results

3.1 The NIS Nonionella sp. T1 oxygen and nitrate respiration and denitrification rates in the Gullmar Fjord

The O_2 respiration rates<u>rate</u> measured <u>infrom</u> the pool of *Nonionella* sp. T1 specimens collected <u>induring</u> the 2018 cruise (GF18) <u>werewas</u> 169 \pm 11 pmol O_2 indiv: d^{-1} with an average BV of 1.3 \pm 0.7 10^{+06} µm³ (BV details, Table 1). The denitrification rate; measured <u>onfrom</u> the same pool of specimens; was 21 ± 9 pmol N indiv: d^{-1} .

The *Nonionella* sp. T1 average BV of the specimens collected induring the 2017 cruise (GF17-3) was $4.0 \pm 0.6 \pm 10^{+06} \, \mu \text{m}^3$, i.e. more than three times larger the *Nonionella* sp. T1 average BV fromof the 2018 eruisesamples $(1.3 \pm 0.7 \pm 10^{+06} \, \mu \text{m}^3)$. As denitrification rates and foraminifera BV are linkedrelated (see method), the measured denitrification rate was corrected using the BV of *Nonionella* sp. T1 from the 2017 cruise. The Hence, the *Nonionella* sp. T1 corrected denitrification rate was $38 \pm 8 \, \mu \text{m}$ pmol N indiv^{*}. $^{-1} \, ^{-1} \, ^{-1} \, ^{-1}$ (Equation S1).

3.2 The NIS Nonionella sp. T1 and foraminifera fauna regarding porewater nitrate micro-distribution

The bottom water at GF17-3 station was oxic (Fig. S1, $[O_2] = 234 \,\mu\text{mol L}^{-1}$) and the measured oxygen penetration depth (OPD) in the sediment was $4.7 \pm 0.2 \,\text{mm}$ (n = 3). No nitrite was revealed on the gel (< 1.7 μ mol L⁻¹), only nitrate was detected. Bottom water average NO₃⁻¹ concentration was $14.6 \pm 2.3 \,\mu\text{mol L}^{-1}$ and nitrate concentration decreased with depth in the sediment (Fig. 5 c, d). Nitrate concentrations ranged between $13.1 \pm 3.2 \,\text{to} \,11.7 \pm 3.4 \,\mu\text{mol L}^{-1}$, from the water-sediment interfaceSWI to the OPD. Nitrate concentrations oncentrations decreased strongly afterunder the OPD from $11.7 \pm 3.4 \,\mu\text{mol L}^{-1}$, from the water-sediment interfaceSWI to the OPD. Nitrate concentrations oncentrations decreased strongly afterunder the OPD from $11.7 \pm 3.4 \,\mu\text{mol L}^{-1}$ untilat 4.0 cm depth. From 4.0 to 5.0 cm depth, NO₃⁻¹ concentration was very low with an average value of $2.7 \pm 0.9 \,\mu\text{mol L}^{-1}$ (Fig. 5 c, d). The PROFILE parameters (Berg et al., 1998) used on laterally averaged nitrate porewater vertical distribution of both stations are available in Table S1. Thus, the PROFILE modelling of the averaged nitrate porewater profiles profile revealed one nitrification zone from 0 to 1.2 cm depth and two denitrifying zones (red line, Fig. 5 d). The first denitrification zone occurred between 1.2 to 3.65 cm depth with a nitrate consumption of 3.92 E^{.05} nmol cm⁻³ s⁻¹ and the second smaller consumption zone was from 3.65 to 5 cm depth (1.53 E^{.06} nmol cm⁻³ s⁻¹). The total denitrification rate from 1.2 to 5 cm depth was 4.07 E^{.05} nmol cm⁻³ s⁻¹ (Fig. 5 d).

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The total densities of living foraminifera were similar between the cores GF17-3A and 3C (Ø 8.2 cm, 5 cm depth) with 1256 individuals and 1428 individuals, respectively (Fig. 5 a and b; Table S2, GF17-3A and 3C). *Nonionella* sp. T1 was the main denitrifying species, accounting for 34 % of the total living fauna in the core GF17-3A and 74 % in GF17-3C (Fig. 5 a, b; Table S3). One other candidate tofor denitrification, *Stainforthia fusiformis*, was in minority: 1 % of the total fauna in both cores (Fig. 5 a, b; Table S3, GF17-3A and 3C). The other known denitrifying species previously reported in the Gullmar Fjord, *Globobulimina turgida* (Risgaard-Petersen et al., 2006) and *Globobulimina auriculata* (Whoele et al., 2018) were absent. Three non-denitrifying species (Piña-Ochoa et al., 2010; Xu et al., 2017; Glock et al., 2019) were dominant in the cores GF17-3A and 3C: *Bulimina marginata* (37 and 5 %, respectively), *Cassidulina laevigata* (9 and 5 %) and *Leptohalysis scotti* (11 and 9 %).

The density and the micro-distribution of *Nonionella* sp. T1 differed between the two cores (Fig. 5 a and b; Table S2, GF17-3A and 3C). In the core GF17-3A and 3C respectively, *Nonionella* sp. T1 density showed large variability from the water-sediment interface to 1.2 cm depth (Table S2) where *Nonionella* sp. Donionella sp. T1 relative abundance accounted for 18 % and 50 % of the fauna in the nitrification zone (Fable S3; from the SWI to 1.2 cm depth) for the cores GF17-3A and 3C; respectively (Table S3). In the firstmain denitrifying zone (from 1.2 cm to 3.65 cm), the *Nonionella* sp. T1 relative abundance represented 27 % and 78 % of the fauna; for the core GF17-3A and 78 % for the core GF17-3C. In the second denitrifying zone, the *Nonionella* sp. T1 relative abundance increased from 3.65 to 5 cm depth and dominated the fauna bywith relative abundances of 60 % and 98%.% (GF17-3A and 3C respectively). The relative abundance of the denitrifying candidate, *Stainforthia fusiformis*, was a minor component in each zones of both cores and did not exceed 2 % (Table S3, GF17-3A and 3C). The three non-denitrifying species (e.g. *B. marginata*, *C. laevigata* and *L. scotti*) also dominated the fauna of both cores GF17-3A and 3C (Table S2 and S4). From the water-sediment interfaceSWI to 1.2 cm depth, *B. marginata* accounted for 42 % and 12 %, *C. laevigata* 16 % and 13 % and *L. scotti* 6 % and 11 %, for the cores GF17-3A and 3C respectively. In the first denitrifying zone (1.2-3.65 cm depth)), *B. marginata* accounted for 34 % and 2 %, *C. laevigata* for 7 % and 2% and *L. scotti* for 25 % and 13 %-% (GF17-3A and 3C respectively-). In the second denitrifying zone (3.65-5 cm depth)), *B. marginata* accounted for 34 % and 0 %, *C. laevigata* was absent and *L. scotti* 5 % and 1 %-% (GF17-3A and 3C respectively-).

Due to severe hypoxia at the GF17-1 station, oxygen was assumed to be below detection limit within the sediment. No nitrite was detected at this station (< 1.7 μmol L⁻¹). Average NO₃⁻ concentration in the bottom water reached 5.7 ± 1.0 μmol L⁻¹ (Fig. 5 g and h). Nitrate concentrations decreased from the sediment surface SWI (4.2 ± 1.0 μmol L⁻¹) to 1.6 cm depth (1.8 ± 0.6 μmol L⁻¹) and then average nitrate concentration remained below the detection limit (1.7 μmol L⁻¹). However, a patchmicro-environment with higher nitrate concentration was visible on the left part of the gel between 2.0 and 3.0 cm depth. A 1D vertical profile passing through this patchmicro-environment (white line, Fig. 5 g) was extracted from the 2D image and the maximal nitrate concentration of thethis patch was above the detection limit with a value of 6.5 μmol L⁻¹ at 2.3 cm depth (blue squaressquare profile, Fig. 5 h). The PROFILE modelling (Table S1) of the laterally averaged nitrate vertical distribution revealed at the sampling time one denitrifying zone from the surfaceSWI to 1.6 cm depth with a nitrate consumption of 2.71 E⁻⁰⁵ nmol cm⁻³ s⁻¹ (red line, Fig. 5 h). BelowNo PROFILE modelling was done under 1.6 cm depth, because nitrate concentration was below the detection limit (hatched grey zone, Fig. 5 h), thus no PROFILE modelling was done after this depth.5 h).

Living foraminifera showed different total densities and a large difference in both species distribution and total densities between the two cores GF17-1A and 1C (Fig. 5 e, f; Table S2); with 1457 individuals and 786 individuals; respectively (Ø 8.2, 5 cm depth). Nonionella sp. T1 represented a low relative abundance of the total fauna with 5 % infor the core GF17-1A and was almost absent (1 %) infor the core GF17-1 C (Table S3). The known denitrifying *G. auriculata* was minor in the fauna with relative abundances of 1 % and 2%; (GF17-1A and 1C respectively). The denitrifying candidate *S. fusiformis* was also found in theboth cores GF17-1A and 1C reaching only 3% of the total fauna (Figure 5 e, f; Table S3). The other denitrifying candidate *B. pseudopunctata*, was almost absent of the total fauna with relative abundances of 0 % and 2 % for the cores GF17-1A and GF17-1C respectively (Table S3). The same three non-denitrifying species as for the observed in oxic station were also dominant infor both cores GF17-1A and 1C: *B. marginata* (64 and 30 %), *C. laevigata* (16 and 15 %) and *L. scotti* (4 and 36 %).

-In the denitrifying zone (0-1.6 cm), Nonionella sp. T1 relative abundance was low, with rare (2 % in%) for the core GF17-1A) and was almost absent from the fauna infor the core GF17-1C. InFor the core GF17-1A, Nonionella sp. T1 relative abundance reached 26 % of the fauna between 1.46 and 2.5 cm depth (Fig. 5 e, GF17-1A), whereas it was almost absent from

the rest of the core GF17-1A and it_was absent from the core GF17-1C (Table S3). InFor the cores GF17-1A and 1C, S. fusiformis reached respectively 2 % and 3 % in the denitrifying zone (0-1.6 cm). InUnder the rest of the cores from denitrifying zone (1.6+to-5 cm-depth,), S. fusiformis represented 4 and 1 % of the fauna; (GF17-1A and 1C respectively.-). The three other non-denitrifying species dominated both cores GF17-1A and 1C. Inin the denitrifying zone (0-1.6 cm-depth)): B. marginata accounted for 66 % and 35 %, C. laevigata 19 % and 19 % and L. scotti 4 % and 24 %-% for the cores GF17-1A and 1C respectively. From 1.6 to 5 cm depth, B. marginata dominated the fauna byaccounted for 61 % and 11 %, C. laevigatalaevigatafor 5 % and 2 % and L. scotti for 6 % and 75 %-% (GF17-1A and 1C respectively.).

4 Discussion

4.1 The NIS Nonionella sp. T1 density in comparison with other species from the Gullmar Fjord

The presence and relative abundance of the NIS Nonionella sp. T1 in the Gullmar Fjord and in the Skagerrak-Kattegat strait hashave been documented during the last decades. The earliest SEM observations of specimens resembling Nonionella sp. T1 morphotype in the deepest part of the fjord date back to summer 1993 (identified as Nonionella turgida, Gustafsson and Nordberg, 2001). The invasive characteristics of Nonionella stella waswere firstly revealed by Polovodova Asteman and Schönfeld, (2015). Then, Nonionella stella was identified as Nonionella sp. T1 morphotype also described as a NIS and potentially invasive species in the Oslofjord by Deldicq et al. (2019). The estimated introduction date of Nonionella sp. T1 into the deepest part of the Gullmar Fjord is 1985 according to Polovodova Asteman and Schönfeld, (2015). The relative abundance of Nonionella sp. T1 in the deepest fjord station was less than 5 % between 1985 and 2007 (Polovodova Asteman and Schönfeld, 2015 and references within). At the GF17-1 hypoxic station, the Nonionella sp. T1 relative abundance was between 1-5 % (Table S3, GF17-1A and 1C). Thus, the Nonionella sp. T1 relative abundance in the deepest part of the fjord seems to remain stable. In contrast to GF17-1 station-GF17-1, the GF17-3 oxic station was sampled for the first time in this study. In this station closer to the mouth of the fjord than GF17-1, the relative abundance of Nonionella sp. T1 varied between 34 and 74 % (Table S3, GF17-3A and 3C). Previous studies showed an increase in the relative abundance of Nonionella sp. T1 represented 10 % of the fauna in June 2013 (Polovodova Asteman and Schönfeld, 2015). The Öresund strait linking the North Skagerrak,

the Kattegat and the Baltic Sea₇ showed an increase in *Nonionella* sp. T1 relative abundance from 1 % to 14 % observed between 1998 and 2009 (Charrieau et al., 2019). The foraminifera fauna in the Gullmar Fjord has changed over the last decennium and *Nonionella* sp. T1 seemedseems to become an invasive species in the Gullmar Fjord oxic shallow water area.

The foraminifera fauna found at the GF17-1 station in the deepest part of the fjord differed from previous studies (Nordberg et al., 2000; Filipsson and Nordberg, 2004; Risgaard-Petersen et al., 2006; Polovodova Asteman and Nordberg, 2013; Polovodova Asteman and Schönfeld, 2015). Indeed, until the early 1980s, the foraminifera fauna in the deepest part of the fjord was dominated by a typical Skagerrak – Kattegat fauna (*Bulimina marginata, Cassidulina laevigata, Hyalinea balthica, Liebusella goësi, Nonionellina labradorica* and *Textularia earlandi*) (Nordberg et al., 2000). However, the fauna changed. *S. fusiformis* and *B. pseudopunctata* became the major species (Nordberg et al., 2000; Filipsson and Nordberg, 2004). Further studies by Polovodova Asteman and Nordberg, (2013) demonstrated that at least until 2011 *S. fusiformis, B. pseudopunctata* and *T. earlandi* dominated the fauna. Foraminifera fauna described in the present study differs it is the

consequence of the occurrence of numerous severe hypoxic events in the fjord (Fig. 3) due to lack of deep-water exchange.

In November 2017 *S. fusiformis* did not exceed 3 % of the fauna (Table S3, GF17-1A and 1C), *B. pseudopunctata* reached only 2 % in the core GF17-1C (Table S3, GF17-1C) and *T. earlandi* was a minor species < 1 %. -Then, in November 2017₂ *B. marginata, C. laevigata* and *L. scotti* were the dominant species in the fjord. The *Elphidium clavatum-selseyensis* species complex (following the definition from Charrieau et al., 2018), *H. baltica, N. labradorica*, and *T. earlandi* were present inwith a low relative abundance (< 5 %, Table S3). Namely, *G. turgida* reached 37 % of the foraminifera fauna in August 2005 at the deepest station (Risgaard-Petersen et al., 2006); whereas in November 2017 this species was minor. The decreasing in relative abundance decreased to become a minor species of the assemblage. However, such trend for *S. fusiformis* and *B. pseudopunctata* must be interpreted with caution since our study used the > 100 μm fraction whereas some of the previous studies used the > 63 μm fraction. We also wet picked the specimens and used Cell Tracker Green to identify living foraminifera, which might affect the results compared to Rose Bengal studies of dry sediment residuals.

The relative abundance of the potential invasive *Nonionella* sp. <u>T1 has increased accordingT1 in 2017increased compared</u> to the study of Polovodova Asteman and Schönfeld₇ (2015) in the oxic part of the fjord. <u>Thelt is also noteworthy that the</u> two non-denitrifying species *B. marginata* and *C. laevigata* described as typical species of the Skagerrak-Kattegat fauna (Filipsson

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and Nordberg, 2004) have again increased markedly in the fjord as well. It is evident that the foraminifera fauna in the Gullmar Fjord is presently very dynamic with considerable species composition shifts, probably following seasonal water body stratification and consecutive oxygen depletion occurring in the fjord (Fig. 3).

4.2 Foraminifera ecology considering porewater nitrate micro-distribution

oxygen lat the rate of 169 ± 11 pmol O₂ indiv⁻¹ d⁻¹ Deeper in the hypoxic zone containing (Fig. 5 c and d). Below the oxygen penetration depth (From 4.7 ± 0.2 mm to 3.5 cm), *Nonionella* sp. T1 canwould store and respires the ambient nitrate (Fig. 5 c, d; from OPD to 3.6 cm depth); *Nonionella* sp. T1 accumulates intracellular nitrate rate to an upper zone where nitrate is still present in the sedimental present in the sedimental present in the sedimental presence to one spatial variability.

Hypoxia occurred approximately at least one month before the sampling cruise in the deepest part of the fjord (Fig. 3). When hypoxia is extended to the water column, nitrification both in the water column and the sediment is reduced or even stopped, as oxygen is almost absent (Fig. 1 b; Childs et al., 2002; Kemp et al., 2005; Conley et al., 2007; Jäntti and Hietanen,

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2012). Under this condition, the coupled nitrification-denitrification processes are strongly reduced (Kemp et al., 1990). At the GF17-1 station, no nitrification in superficial sediment was showed by our data and nitrate was low but still detectable in the bottom water. Nitrate can diffuse from the water column into the sediment, and thereby generate the denitrification zone as modelled by PROFILE between the surfaceSWI and 1.6 cm depth (Fig. 5 h).

The rare presence of the NIS Nonionella sp. T1 and other denitrifying species as Globobulimina auriculata, Bolivina pseudopunctata and Stainforthia fusiformis in the hypoxic station indicates that sediment chemical conditions turned unfavorable towards denitrification during prolonged hypoxia. Instead, the non-denitrifying species Bulimina marginata, Cassidulina laevigata, and Leptohalysis scotti dominated in this hypoxic environment. Their survival could be due to seasonal dormancy (Ross and Hallock, 2016; LeKieffre et al., 2017) and their ability to release propagules which can disperse and reproducegrow when environmental conditions turn favorable again (Alve and Goldstein, 2003). The suspected deep nitrification zone (blue square profile, Fig. 5 h) could explainindicate the presence of nitrate micro-niches deeper in the sediment and might explain the patchy distribution of Nonionella sp. T1 also at the hypoxic site (see Fig. 5 e; Table S2, GF17-1A). Therefore, deep nitrate production in these micro-environments could favor the presence of Nonionella sp. T1, which can be attracted by this nitrate source of electron acceptor to respire (Nomaki et al., 2015; Koho et al., 2011). This deep nitrification zone could be athe result of an aerobic or anaerobic process. An aerobic nitrification zone in deep sediment can be formed by macrofaunal activity (burrowing activity) that introduce some oxygen deeper into anoxic sediment (Aller, 1982; Karlson et al., 2007; Nizzoli et al., 2007; Stief, 2013; Maire et al., 2016). This nitrification zone could also be due to an anaerobic process. The Gullmar Fjord is Mn-rich (Goldberg et al., 2012) and metal-rich particles can be bio-transported into the anoxic sediment, thus allowing ammonium oxidation into NO3- by Mn and Fe-oxides in the absence of oxygen deeper in the sediment (Aller, 1994; Luther et al., 1997).

4.3 Contributions and potential impacts of the NIS Nonionella sp. T1 to benthic denitrification in the Gullmar Fjord

If we considerConsidering that *Nonionella* sp. T1 is denitrifying the nitrate from sediment porewater (approach A, Table 2; see method 2.5)), its contribution to benthic denitrification in the oxic station would be 47 % in the core GF17-3A and would reach 100 % in the core GF17-3C. If we consider that *Nonionella* sp. T1-also uses its intracellular nitrate pool for

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denitrification (approach B), its contribution to benthic denitrification would be 32 % in the core GF17-3A and would reach 50 % in the core GF17-3C (Table 2). These two calculation approaches highlight the difficulties and the importance of knowing the concentration of environmental nitrate and foraminifera intracellular nitrate at the same time to estimate the contributions of foraminifera to benthic denitrification. Moreover, in this study there is no data on anammox process which contributes also into the total denitrification (Brandes et al., 2007). The results reported in previous studies as Engström et al., (2005) do not allow us to extrapolate their data at our oxic station, located at the entrance of the fjord. Thus, we assume that our estimate of denitrification is conservative; since the possible contribution of anammox is not included in the calculation. However, despite these uncertainties *Nonionella* sp. T1 contribution to benthic denitrification supports the hypothesis that this non-indigenous denitrifying foraminifer playplays a major role in the benthic nitrogen cycle-for sediments.

At the hypoxic station, the opposite was shown where the estimated contribution of *Nonionella* sp. T1 to benthic denitrification was below 1 % whatever the calculation approach. The estimated contributions of the other denitrifying foraminifera found in the hypoxicthis station were low. Foraminifera contributed to almost 5 % of benthic denitrification—in the hypoxic station. Compared to the oxic station, the NIS *Nonionella* sp. T1 and the other denitrifying species contributions to benthic denitrification were smallweak in a prolonged hypoxic station of the Gullmar Fjord.

Overall, the Gullmar Fjord is well oxygenated except for the deepest basin where oxygen goes down when there is no deep water exchange (Fig. 3 c). Therefore, the GF17-3 oxic station could be considered representative of the Gullmar Fjord benthic ecosystem. *Nonionella* sp. T1 is not the most efficient denitrifying species compared to *Globobulimina turgida* (42 pmol N ind⁻¹ d⁻¹, with BV = 1.3 10⁺⁰⁶ µm³) and also less efficient than *Nonionella* cf. *stella* from Perú. However, *Nonionella* sp. T1 high density could accelerate sediment denitrification and participate to increase the contrast between the two hydrographic conditions. Indeed, an increase in contrast due to the increasing discrepancy of bottom water oxygenation conditions; oxic vs severe hypoxia induced between stations induces a gap in the availability of nitrate for anaerobic facultative metabolisms in the sediment. In the oxygenated part of the fjord, high contribution to benthic denitrification (estimated between 50 and 100%) by *Nonionella* sp. T1 could contributetake part to a potential de-eutrophication of the system by increasing the nitrogen loss. Primary production (PP) efin the Gullmar Fjord is dominated by diatoms bloomdiatom blooms in spring and

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autumn (Lindahl and Hernroth, 1983). Since the 1990s, Lindahl et al. (2003) observed an increase in PP ofin the Gullmar Fjord, therefore a potential eutrophication potentially changing its trophic status towards eutrophic. This increase in PP also shown in the adjacent Kattegat could be related to the nitrogen input loading from the land and atmosphere (Carstensen et al., 2003). Lindahl et al. (2003); argued that the PP ofin the Gullmar Fjord was due to climatic forces resulting from a strong positive North Atlantic Oscillation (NAO) index, which increased the availability of deep-water nutrients (Kattegat nitrate-rich) and due to warmer ocean through changes in the thermocline. The benthic denitrification of the Gullmar Fjord produces nitrogen unassimilable by primary producers. Moreover, denitrifying foraminifers foraminiferal nitrate uptake and intracellular nitrate becomes unavailable to the system and can bestorage act as an additional sink through bio-transportedtransportation and permanently sequestered permanent sequestration in sediments (Glock et al., 2013; Prokopenko et al., 2011). Thus, denitrifying foraminifera including Nonionella sp. T1 could help counterbalance a potential eutrophication of the system via nitrogen loss (Seitzinger, 1988).

Whereas Contrastingly, in the hypoxic partspart of the fjord, nitrate and nitrite rapidly exhausted become scarce, resulting in a decrease in benthic denitrification. The consequence is a decrease of denitrifying including foraminifera fauna. The increase of ammonium in anoxic sediment resulting by a decrease incontribution. As a consequence of oxygen and nitrate scarcity, nitrification, denitrification and anammox processes does not allow the are less intense resulting in a decrease of nitrogen elimination from the sediment to the water column. Thus, potentially promoting an ammonium mitigation and accumulation of ammonium in the decrease part of the fjord parts subjected to prolonged severe hypoxia (Fig. 1). Moreover, the low availability of nitrate in the sediment would possibly increase the benthic transfer towards the water column of reduced compounds such as manganese and iron produced deeper in the sedimentary column by other anaerobic metabolisms (Hulth et al., 1999). These new results demonstrate that the role of denitrifying foraminifera is underestimated in the nitrogen cycle and that overlooking this part of the meiofauna may lead to a misunderstanding of environments subject to hydrographic changes.

5 Conclusion

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This study revealed a drastic change in living foraminifera fauna due to several hypoxic events that occurred in the last decennium in the Gullmar Fjord. For the first time, the non-indigenous species (NIS) *Nonionella* sp. T1 dominated up to 74 % the foraminifera fauna at a station with oxygenated bottom waters and high nitrate content in sediment porewater. This NIS can denitrify up to 50-100 % of the nitrate porewater sediment under oxic conditions in the fjord. Whereas, under prolonged hypoxia, nitrate depletion turns environmental conditions unfavorable for foraminifera denitrification, resulting in a low density of *Nonionella* sp. T1 and other denitrifying species. Foraminifera contribution to benthic denitrification was negligible (~ 5 %) during prolonged seasonal hypoxia in the fjord. Moreover, the potential invasive denitrifying *Nonionella* sp. T1 could impact the nitrogen cycle under oxic conditions by increasing the sediment denitrification and could counterbalance potential eutrophication of the Gullmar Fjord. Our study demonstrated that the role of denitrifying foraminifera is underestimated in the nitrogen cycle especially in oxic environments.

FiguresFigure list

(a) Oxic bottom water

(b) Hypoxic bottom water

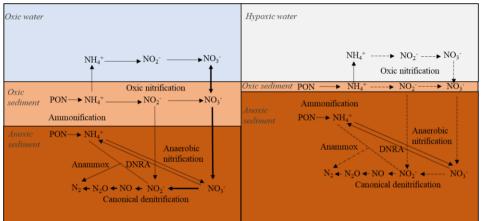


Figure 1. Simplified nitrogen cycling in marine sediments when the bottom water is oxic (a) and hypoxic (b). Chemical formulae: PON (particulate organic nitrogen), NH_4^+ (ammonium), NO_3^- (nitrate), NO_2^- (nitrite), NO_2^- (nitrogen oxide), N_2O (nitrous oxide), N_2 (nitrogen). The bold/dotted arrows indicate reactions advantaged/reduced by oxygen and nitrate presence/depletion. See text for more details. Modified from Jantti and Hietanen, (2012),

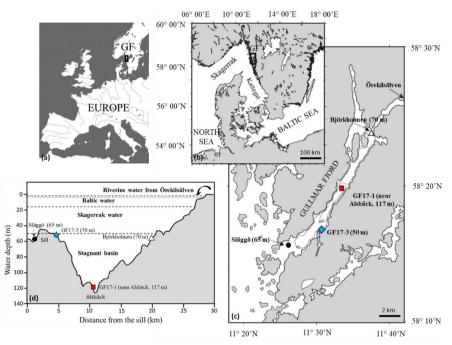
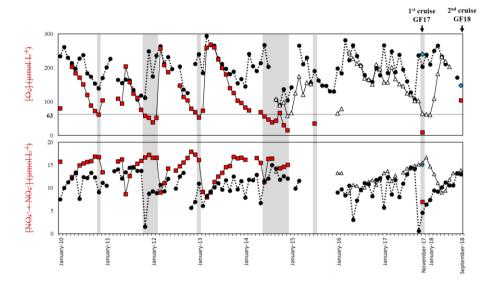


Figure 2. (a-c) Location of studied stations in the Gullmar Fjord (Sweden); blue diamond: GF17-3 oxic station (50 m water depth); red square: GF17-1 hypoxic station (117 m water depth); dark circles: monitoring stations Släggö (65 m water depth) and Björkholmen (70 m water depth). (d) Transect from the sill with the four Gullmar Fjord water masses and the studied stations (modified from Arneborg et al., 2004).



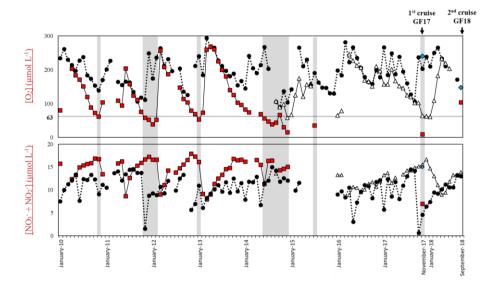


Figure 3. Record from January 2010 to September 2018 of bottom water oxygen ([O₂]) and nitrite + nitrate ([NO₃ $^{-}$ + NO₂ $^{-}$]) measurements from the monitoring stations Släggö (65 m water depth; black dot), Björkholmen (70 m water depth; white triangle) and the sampling stations GF17-1 (Alsbäck, 117 m water depth; red square) and GF17-3 (50 m water depth; blue diamond). The arrows indicate the date of the two sampling cruises: the 2017 cruise (14th, 15th November 2017) and the 2018 cruise (5th September 2018). The grey zones indicate hypoxiahypoxic periods with a threshold (fof [O₂] < 63 μ mol L⁻¹)_{2.}

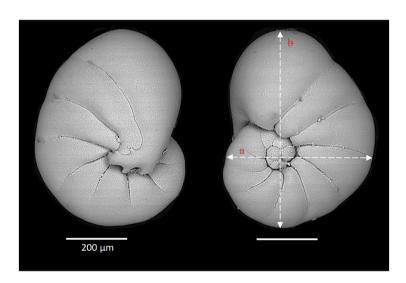


Figure 4. Scanning Electronic Microscope images of a *Nonionella* sp. T1 from the GF17-3 oxic station in the Gullmar Fjord. White lines (a, b) correspond to measured distances serving for a spheroid prolate volume model.

Table 1. Total shell volume (μm^3) and the biovolume (BV, μm^3) corresponding to 75_% of the total shell volume measured on the pool of five *Nonionella* sp. T1 from the 2017 and the 2018 cruises in the Gullmar Fjord. Abbreviations: sd (standard deviation), ind. (individual).

Nonionella sp. T1	1 st cruise total shell volume	1 st cruise BV	2 nd cruise total shell volume	2 nd cruise BV
ind. 1	6.7 10 +06	5.0 10 +06	3.1 10 +06	2.3 10 +06
ind. 2	4.5 10 +06	3.4 10 +06	2.4 10 +06	1.8 10 +06
ind. 3	5.1 10 +06	3.8 10 +06	1.4 10 +06	1.0 10 +06
ind. 4	4.9 10 +06	3.7 10 +06	9.2 10 +05	6.9 10 +05
ind. 5	5.8 10 +06	4.4 10 +06	6.2 10 +05	4.7 10 +05
Average (µm³)	5.4 10 +06	4.0 10 +06	1.7 10 +06	1.3 10 +06
sd (μm³)	0.8 10 +06	0.6 10 +06	1.0 10 +06	0.7 10 +06

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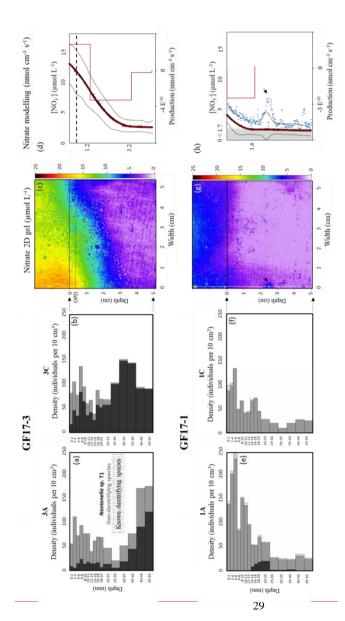
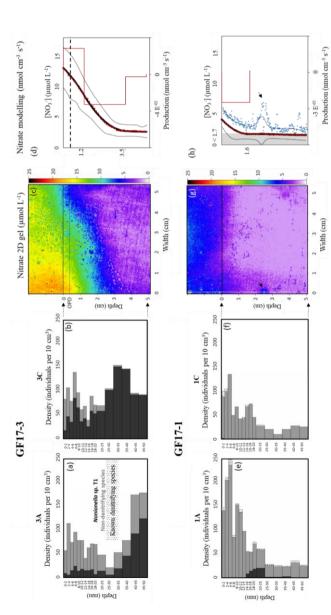


Figure 5. Micro-distributions of living foraminifera densities in GF17-3 oxic station (a, b) and in GF17-1 hypoxic station (e, f), Nonionella sp. T1 specimens are in black, the sum of the non-denitrifying species in grey colors and the small dots (e, f) show the other denitrifying pixel line of the nitrate distribution image (290 pixels wide), the standard deviation is represented by two fine dotted lines (e and g with PROFILE. The 1D profile corresponding to x = 1 mm (white line, g) is represented with a blue square profile (h) and the deep nitrate The sediment-water interface is represented by a black line at 0 cm depth (c, g) and the Oxygen Penetration Depth (OPD) is represented by the dashed line in bold at 4.7 ± 0.2 mm depth (c). Nitrate 1D profiles (d and h, black dots) are calculated using the average value of each respectively). The corresponding best fitting concentration profiles (red dots, d and h) and the production zones (red line) are modelled species (known and potential candidates). The maps of porewater nitrate 2D gels are presented for stations GF17-3 (e) and GF17-1 (g). spot is indicated by a black arrow. The hatched grey zone (h) represents the detection limit of the nitrate $2D \gcd(<1.7 \mu mol L^4)$.



(known and potential candidates). The maps of porewater nitrate 2D gels are presented for the stations GF17-3 (c) and GF17-1 (g). The SWI is represented by a black line at 0 cm depth (c, g) and the OPD is represented by the dashed line in bold at 4.7 ± 0.2 mm depth (c). Figure 5. Micro-distributions of living foraminifera densities in GF17-3 oxic station (a, b) and in GF17-1 hypoxic station (e, f). Nonionella TI specimens are in black, the sum of the non-denitrifying species in grey and the small dots (e, f) show the other denitrifying species pixels wide), the standard deviation is represented by two fine dotted lines. The corresponding best-fitting concentration profiles (red dots, d and h) and the production zones (red line) are modelled with PROFILE software. The 1D profile corresponding to x = 1 mm (white line, g) is represented with a blue square profile (h) and the deep nitrate spot is indicated by a black arrow. The hatched grey zone (h) represents the detection limit of the nitrate 2D gel (<1.7 μ mol L⁻¹). Nitrate 1D profiles (d and h, black dots) are calculated using the average value of each pixel line of the nitrate distribution image (290

Table 2. Summary of the NIS Nonionella sp. T1 contributions to benthic denitrification in the Gullmar Fjord. The porewater denitrifications denitrification, zones come from PROFILE modelling (Fig. 5 d, h). To estimate the contributions of Nonionella sp. T1 the number of counted specimens per zones was used. Two different approaches were used to estimate the contribution of Nonionella sp. T1: (A) divided the Nonionella sp. T1 denitrification rate divided by the nitrate porewater denitrification rate estimated from PROFILE modelling, then the second approach; (B) divided the Nonionella sp. T1 denitrification rate divided by thenitrate porewater, denitrification rate from PROFILE plus the Nonionella sp. T1 denitrification rate. The calculations are detailed in Equation S2.

Stations	Sediment depth interval of	Nonionella sp. T1 (counted	Nitrate porewater denitrification	Nonionella sp. T1 denitrification	Nonionella sp. T1 contribution	
	denitrification (cm)	specimens per zone)	rates (nmol cm ⁻³ s ⁻¹)	rates (nmol cm ⁻³ s ⁻¹)	(%), approach A	(%), approach B
GF17-3A	1.2 to 5	841	4.07 E ⁻⁰⁷	1.90 E ⁻⁰⁵	47	32
GF17-3C	1.2 to 5	1807	4.07 E ⁻⁰⁷	4.06 E ⁻⁰⁵	100	50
GF17-1A	0 to 1.6	3	2.71 E ⁻⁰⁵	6.72 E ⁻⁰⁸	0	0
GF17-1C	0 to 1.6	12	2.71 E ⁻⁰⁵	2.69 E ⁻⁰⁷	1	0

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Author contributions

C.C. participated in the sampling cruise, did the foraminifera taxonomy, contributed to 2D gel experiments and analyses by hyperspectral camera. C.C. did the nitrate and oxygen respiration measurements and wrote the present manuscript. E.G. participated in the sampling cruise, contributed to foraminifera analysis, scientific discussions. E.M. participated in the sampling cruise, managed with A.M. the 2D gels experiments, and contributed to hyperspectral camera treatments and scientific discussions and manuscript rewriting. H.L.F managed with A.M the sampling cruise and contributed to foraminifera taxonomy and scientific discussions and manuscript rewriting. N.R.P. managed the oxygen and nitrate respiration measurements and contributed to the scientific discussions. P.L. managed hyperspectral treatments for 2D gels and contributed to scientific discussion. M.G. participated in the 2D gel lab experiments and hyperspectral treatments. T.J. participated to the sampling cruise, contributed to 2D gels experiments and scientific discussions and manuscript rewriting. B.J. contributed to hyperspectral camera treatments and scientific discussions and manuscript rewriting.

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Competing interests

The authors declare no competing interest.

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