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Contribution of ammonium oxidizing archaea and bacteria to intensive
nitrification during summer in Mutsu Bay, Japan.

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Abstract

Nitrification, in which ammonium is oxidized to nitrate via nitrite, is a regeneration process of bioavailable nitrogen. Ammonia-oxidizing archaea (AOA) and ammonia-oxidizing bacteria (AOB) carry out the ammonium oxidation. Here we measured the ammonium oxidation rates (AORs), distinguishing those of AOA (V_A) and AOB (V_B), for two years in Mutsu Bay, a subarctic semi-enclosed bay. Nitrate and ammonium were depleted in the upper layer in summer, but a high concentration of nitrate was observed at the bottom (below 40 m), of up to $7.8 \mu\text{M}$ with a strong vertical gradient within several meters of the seafloor. A High AOR of up to $0.17 \mu\text{M d}^{-1}$ were observed in the bottom waters. The AOA were responsible for most of the total AOR, but the V_B exceeded the V_A just above the seafloor (0.3 m) in September. Removal of the suspended solids (SS) did not change the AOA's oxidation activity but eliminated the AOB's. This finding suggested that AOA were free-living whereas AOB were attached to the SS's surface. It is also suggested that the sediment was this SS's source. When ammonia mono-oxygenase, *amoA*, of AOA and AOB, was quantified, the copy number of AOA-*amoA* was an order of magnitude higher than that of AOB-*amoA* throughout the study period. A significant correlation ($p < 0.05$) was found between the AOB-*amoA* copy number and V_B whereas no correlation was found between the AOA-*amoA* copy number and V_A , suggesting the AOA-*amoA* copy number did not always represent active cells. Ammonia oxidation was occurring near the bottom waters of Mutsu Bay during the summer, and it was accompanied by a dynamic balance reflecting the difference in the AOA's and AOB's lifestyles, free-living and particle-associated, respectively.

1 Keywords: Nitrification, oligotrophic bay, AOB, AOA, *amoA*

2

1. Introduction

Phytoplankton utilize three forms of inorganic nitrogen (N) species to support their growth: nitrate (NO_3), nitrite (NO_2), and ammonium (NH_4). These N species are supplied to the euphotic layer from the terrestrial, deep water; sedimentary sources; and decomposition of detrital organic matter. During the remineralization process of organic N, NH_4 is the first product and is further oxidized to NO_3 . This process is called nitrification. Nitrification consists of the two microbially mediated processes by which NH_4 is oxidized through NO_2 to NO_3 . Light inhibits nitrification because NH_4 - and NO_2 -oxidizing organisms are sensitive to light (Olson, 1981). Therefore, light suppresses the production of NO_3 by nitrification in the euphotic layer. This finding indicates that NO_3 is produced below the euphotic layer and supplied through advection and mixing into the euphotic layer (Grunde and Juniper, 2011).

Marine nitrifying organisms comprise ammonium-oxidizing bacteria (AOB), ammonium-oxidizing archaea (AOA), and nitrite-oxidizing bacteria (NOB)(Table 1). Archaea have been found to be responsible for a significant proportion of NH_4 oxidation (Könneke et al., 2005; Wuchter et al., 2006). Species composition and distribution of AOA and AOB have been studied using PCR-based techniques (Caffrey et al., 2003; O'Mullan and Ward, 2005; Pedneault et al., 2014). In most studies to characterize and quantify AOA and AOB, researchers have used the functional and phylogenetic marker gene *amoA* (ammonia mono-oxygenase subunit A). Several studies have shown that the abundance of AOA is a few orders of magnitude higher than that of AOB (Bouskill et al., 2012; Newell et al., 2011; Ward and O'Mullan, 2002; Wuchter et al., 2006).

It has been reported that the ammonia oxidizers' abundance, rather than their community structures, could explain the changes in nitrification rates (Hou et al, 2013).

1 However, AOA dominate in abundance, but AOB drive gross nitrification in land soil
2 (Jia and Conrad, 2009, Sterngren et al. 2015). Therefore, concerns have arisen about not
3 only the relationship between *amoA* gene abundance and potential nitrification rate
4 (PNR) but also AOA's and AOB's ecological importance in nitrification (Ando et al.,
5 2009; Baker et al., 2012; Bernhard et al., 2007). The PNR was measured ammonium
6 oxidation rate (AOR) at a higher NH_4 concentration (more than ten times) than in situ
7 concentration whereas AOR was measured at the same order of magnitude of the in situ
8 concentration. The PNR was significantly correlated with archaeal *amoA* gene
9 abundance but had no correlation with bacterial *amoA* gene abundance in the Yangtze
10 Estuary, indicating AOA's importance (Zheng et al., 2014). In a few studies, researchers
11 have described the composition of ammonium-oxidizing organisms (AOOs) in
12 conjunction with ammonium oxidation activity at the in situ concentration level (AOR)
13 in the marine systems.

14 Mutsu Bay is a semi-enclosed bay, located at the northern end of Honshu, Japan
15 (Fig. 1). Its area and average depth were 1,667 km^2 and 38 m. The aquaculture of
16 scallop (*Patinopecten yessoensis*) and other fishing activities have been performed for a
17 long time in this bay. The bottom topography is flat, deepening gradually toward the
18 mouth. Seawater exchange occurs through the narrow mouth (opening width, 10 km)
19 with Tsugaru warm water. A decreasing trend of nutrient concentration
20 (Oligotrophication) has been discovered in the last three decades with the long-term
21 monitoring of nutrients in the bay (Kudo et al. 2014). Dissolved inorganic nitrogen
22 (DIN: $\text{NO}_3 + \text{NO}_2 + \text{NH}_4$) decreased to one-third its original amount at 20 m and two-
23 thirds at the bottom from the 1980s to the 2010s. This oligotrophication was thought to
24 occur because the introduction of scallop aquaculture induced the acceleration of faces

origin organic matter deposition to the sediment. N seems to be a limiting nutrient of primary productivity in the bay because the N:P ratio (~10) is lower than the Redfield ratio (i.e., 16). Therefore, the nitrification process is the most important process in primary productivity in the bay. Ammonium oxidization through the microbial process is a key factor to evaluate the possibility of sustainable use for fishing grounds because ammonium oxidation is the rate-limiting step in nitrification. Further knowledge of AOA's and AOB's role in nitrification is needed in the enclosed bay environment receiving organic matter from human and aquaculture activity. We hypothesize that the role and contribution of AOA and AOB may vary responding to the environmental change in the bottom layer explaining the seasonal variation in AOR. The objectives of this study were to elucidate the seasonal variation in AOR and AOA's and AOB's relative contribution to AOR in the bottom layer of Mutsu Bay. The relative contribution was evaluated by AOR using an inhibitor of ammonium oxidation by AOB (Bianchi et al., 1997; Painter, 2011). The *amoA* gene was quantified by quantitative PCR (qPCR) targeting for AOB and AOA. The contribution of particle-associated organisms to ammonium oxidation near the bottom was also examined.

2. Materials and Method

2.1 Sampling

Water samples were collected during seven cruises in May, early September, late September, and November 2018 and July, September, and November 2019 by T/S *Ushio Maru* at N-43 (water depth: 50 m), which is located in the eastern part of Mutsu Bay (Fig. 1). Salinity, temperature, and dissolved oxygen (DO) were monitored with a Sea-Bird 19 plus CTD (Conductivity, Temperature, and Depth) sensor. Water samples

were taken from the layers 2 m, 1 m, and 0.3 m above the seafloor (hereafter referred to as B-2 m, B-1 m, and B-0.3 m, respectively). Water sampling at B-2 m and B-1 m was conducted with a horizontal X-Niskin sampler. Water near the seafloor was sampled with a triple-tube mounted undisturbed corer (Type-GS, Rigo, Tokyo, Japan) and siphoned from the overlying water of the undisturbed sediment cores. This water was a mixture of the overlying water 0 cm-30 cm above the sediment and labeled B-0.3 m.

Seawater samples for nutrient analysis were collected and stored frozen until analysis. An aliquot of sample seawater was filtered with a pre-weighed GF/F (Whatman, 25 mm diameter, pre-treated at 450 °C for 6 h) for suspended-solids (SS) measurement. The filter was stored frozen, freeze-dried, and weighed several times until a constant value was attained.

2.2 Ammonium oxidation rate (AOR) by AOA and AOB

The bottom water was transferred to a light-shaded 60-mL syringe. The sample was amended with ammonium sulphate ($(\text{NH}_4)_2\text{SO}_4$) at 3-4 $\mu\text{mol N L}^{-1}$. Sodium chlorite (NaClO_3), an inhibitor of nitrite oxidation, was added at a final concentration of 10 mmol L^{-1} using a three-way stopcock to prevent its exposure to air. The samples for measuring AOR by AOA were supplemented with Allyl-thiourea (ATU), an inhibitor of ammonium oxidation by AOB, at a final concentration of 86 $\mu\text{mol L}^{-1}$ (=10 mg L^{-1}). ATU at 86 $\mu\text{mol L}^{-1}$ inhibits AOB but not AOA (Ginestet et al, 1998; Santoro et al, 2010). Therefore, the samples for measuring total AOR by AOA and AOB were not added to ATU. Three replicates were prepared for each condition. Incubation was conducted at the in situ temperature in darkness for 74 h. During the incubation, subsamples for nutrient analysis were taken four to five times (0, 24, 36, 48, and 72 h of

incubation) from the syringes into sterile screw spits and frozen until analysis. The control was prepared using sterilized seawater with the addition of all the chemicals mentioned above.

After the samples were thawed, and nutrients (NH_4^+ , NO_2^- , NO_3^-) were analyzed using a QuAatro (Bran-Luebbe, Tokyo, Japan) following Grasshoff et al. (1999).

The total AOR by AOA and AOB (V_{A+B}) was obtained by the slope of the time-course change in NO_2 concentration for each syringe without ATU. The AOR by AOA alone (V_A) was obtained by the slope of the time-course change in NO_2 concentration where the AOB ammonia oxidation inhibitor ATU was added. The difference between the two rates ($V_{A+B}-V_A$) was defined as the AOR by AOB (V_B). If the average slope of triplicate samples was not significantly different from the control (Mann-Whitney U-test, $p > 0.05$), then AOR was denoted as nd (not detected).

To evaluate the contribution of SS to AOR, an inline filtration through a 3- μm pore-size filter (Nuclepore) was conducted for the B-0.3 m waters in September 2019. The AOR was measured for the filtrate as described above.

2.3 DNA purification from filter

Subsamples were taken for gene analysis of AOA and AOB in 2018 directly from the outlet of the sampler connected to a 60-mL sterilized syringe and filtered through a Sterivex filter (Millipore, 0.2- μm pores) and stored at -80 °C. Subsamples were also taken during the incubation to measure AOR.

After the sample was thawed, the filter was removed from the Sterivex cassette and DNA was extracted following the Campbell method (Wilson, 2001). The filters were chopped into 2-mL tubes, and CTAB reagent (2.0 % CTAB, 1.0 % PVP, 1.4 M

NaCl, 20 mmol L⁻¹ EDTA, 100 mmol L⁻¹ Tris-HCl, pH 8.0) was added, followed by incubation at 65°C for 15 min and then incubation with chloroform:isoamyl alcohol = 24:1 (CI). The tubes were stirred at room temperature for 20 min and then centrifuged at 4 °C (12,500 rpm, 15 min). CI was added to the supernatant and centrifuged at 4 °C (12,500 rpm, 15 min). The supernatant was mixed with 0.3 M sodium acetate (pH 5.2); then an equal volume of isopropanol was added. This mixture was cooled at -80 °C for 2 h to precipitate DNA. DNA was precipitated by centrifugation at room temperature (13,000 rpm, 30 minutes), and the supernatant was removed by decantation. The DNA was washed with 70% ethanol and centrifuged (13,000 rpm, 5 min) at room temperature. The pelleted DNA was vacuum dried and dissolved in 50 µL of TE buffer (10 mmol L⁻¹ Tris-HCl, 1 mmol L⁻¹ EDTA, pH 8.0).

2.4 Quantitative PCR (qPCR) for *amoA*

The CFX96 Real-Time system (BioRad Laboratories) was used for qPCR analysis. The reaction solution consisted of 10 µL of SsoFast EvaGreen Supermix (BioRad), 1 µL of each primer (final concentration 500 nmol L⁻¹), 1 µL of sample DNA (1-5 ng), and 7 µL of sterile water to make the final volume 20 µL. The primers used were crenAMO_F and crenAMO_R for Archaeal *amoA* (AOA- *amoA*) (Hallam et al., 2006) and AmoA-1F and AmoA-2R for bacterial *amoA* (AOB- *amoA*) (Rotthauwe et al., 1997) (Table 2). The qPCR was programmed to denature and activate polymerase at 95 °C for 30 s for the first cycle and programed at 95 °C for 5 s, annealing at 55 °C for 20 s (AOA-*amoA*) or 57 °C for 20 s (AOB-*amoA*), extension at 72 °C for 20 s for the second cycle. This program was performed for 39 cycles. The fluorescence for melting curves was measured in the range of 60-95 °C with a 0.5 °C increase every 5 s. Each

sample was measured three times and normalized as the number of copies per mL of test water. The confidence range of the calibration curve was 3.1×10^6 - 3.1×10 copies mL^{-1} for AOA- *amoA* and 3.8×10^6 - 3.8×10 copies mL^{-1} for AOB- *amoA*. For the standard curve, the PCR product for AOA-*amoA* and AOB-*amoA* from test water was cloned with p-GEM-T Easy vector (Promega) by TA cloning. The cloned plasmid was transformed to *E. coli* DH5 α and purified with a Qiagen plasmid purification Mini kit (QIAGEN). These plasmids were used for qPCR standard templates.

3. Results

3.1 Physical and nutrient environments

The vertical profile of temperature (Fig. 2) indicated that the thermocline developed in July and September. The temperature in the bottom layer (> 40 m) was below 15°C from May to July, and it was between 16 and 22°C in September. The temperature in November was vertically uniform at 17°C in 2018 and 2019.

DIN varied from 0.02 to $1.93\ \mu\text{M}$ for NH_4 , from 0.01 to $1.68\ \mu\text{M}$ for NO_2 , and from 0.02 to $7.84\ \mu\text{M}$ for NO_3 (Fig. 3). DIN in the shallower depths (< 30 m) was low (below $1\ \mu\text{M}$). DIN concentrations in the bottom layer were highest in September. The highest NH_4 and NO_2 concentrations throughout the study period were found in the bottom layer on July 27 and September 26, 2019, at $1.9\ \mu\text{M}$ and $1.7\ \mu\text{M}$, respectively. The highest NO_3 concentration was found in the bottom layer on September 23, 2018, at $7.8\ \mu\text{M}$. In contrast, DIN concentrations in May and November were mostly below $2\ \mu\text{M}$.

3.2 Ammonia oxidation rate (AOR) and contribution of AOA and AOB

1 The total AOR (V_{A+B}) at B-1 and B-2 m varied from nd to $0.10 \mu\text{M d}^{-1}$ from
2 early September 2018 to November 2019 (Fig. 4). The V_{A+B} at B-0.3 m varied from
3 0.007 to $0.173 \mu\text{M d}^{-1}$ from May 2018 to November 2019. The V_{A+B} was generally high
4 in September. The maximum V_{A+B} was observed at B-0.3 m in late September 2018.

5 The AORs by AOA (V_A) and AOB (V_B) in the bottom layers varied from 0.04 to
6 $0.104 \mu\text{M d}^{-1}$ and from nd to $0.099 \mu\text{M d}^{-1}$ from May 2018 to November 2019,
7 respectively (Fig. 5). The V_A tended to have a certain range, but V_B showed the
8 increasing trend to the seafloor.

9 AOA's and AOB's contribution to the total AOR in September was different at
10 B-2 m, B-1 m, and B-0.3 m (Fig. 6). AOA's and AOB's average contribution was 86%
11 and 14% at B-2 m, 57% and 43% at B-1 m, and 41% and 59% at B-0.3 m, respectively.
12 Therefore, AOA contributed more than AOB at B-1 m, and AOB contributed more at
13 B-0.3 m.

14 15 3.3 Contribution of suspended solids (SSs) to AOR

16 The concentration of SSs at B-0.3 m in September 2019 was $135 \pm 13 \text{ mg L}^{-1}$.
17 After filtration with a $3\text{-}\mu\text{m}$ filter, it dropped to $20 \pm 15 \text{ mg L}^{-1}$ (Fig. 7a). The filtration
18 eliminated 85% of SSs.

19 The total AOR (V_{A+B}) in the unfiltered water at B-0.3 m was $0.11 \pm 0.037 \mu\text{M d}^{-1}$.
20 ¹. The V_{A+B} in the filtered water was $0.050 \pm 0.019 \mu\text{M d}^{-1}$ in September 2019 (Fig. 7b).
21 A 55% decrease in the V_{A+B} was observed after filtration (t-test, $p < 0.05$). The
22 contribution of AOB accounted for 66% of the total AOR in the unfiltered B-0.3 m
23 water, but it was almost zero in the filtered water. However, there was no significant
24 difference in the V_A between the filtered and the unfiltered samples ($p > 0.05$).

3.4 Spatiotemporal variation of AOA and AOB copy number

The AOA-*amoA* copy number varied from nd to 3.6×10^4 copies mL⁻¹ at B-2 m, $0.034\text{--}6.3 \times 10^4$ copies mL⁻¹ at B-1 m, and $1.1\text{--}130 \times 10^4$ copies mL⁻¹ at B-0.3 m from May to November 2018 whereas that of AOB-*amoA* varied between $0.017\text{--}15 \times 10^2$ copies mL⁻¹ at B-2 m, $0.0045\text{--}2.3 \times 10^3$ copies mL⁻¹ at B-1 m, and $0.36\text{--}6.6 \times 10^2$ copies mL⁻¹ at B-0.3 m during the same period (Fig. 8). The AOA- *amoA* and AOB- *amoA* copy numbers were larger in September in all samples except AOA- *amoA* at B-0.3 m in May. The AOB-*amoA* copy numbers were larger at B-1 m and B-2 m than at B-0.3 m. The AOA-*amoA* was 10-100 times higher than the AOB-*amoA* in all samples except at B-2 m in May 2018, when AOA-*amoA* was ns.

The AOA-*amoA* copy number after 24 h of incubation was $0.22\text{--}0.62 \times 10^4$ copies mL⁻¹ at B-2 m, $0.58\text{--}1.89 \times 10^5$ copies mL⁻¹ at B-1 m, and $0.01\text{--}2.39 \times 10^5$ copies mL⁻¹ at B-0.3 m. The AOB-*amoA* copy number after 24 h of incubation was $0.16\text{--}1.82 \times 10^3$ copies mL⁻¹ at B-2 m, $0.37\text{--}2.18 \times 10^3$ copies mL⁻¹ at B-1 m, and $0.037\text{--}5.96 \times 10^3$ copies mL⁻¹ at B-0.3 m. A significant correlation ($p < 0.05$) was found between the AOB-*amoA* copy number after 24 h and V_B whereas no correlation was found between the AOA-*amoA* copy number and V_A (Fig. 9).

4. Discussion

4.1 Temporal variation of temperature and nutrients

In summer, a thermocline was observed in the bottom water. In November, the thermocline disappeared due to winter vertical mixing and the water temperature became uniform in the water column. The concentrations of NO₂ and NO₃ in the bottom

water during the observation period were highest in September in 2018 and 2019 (Fig. 3). The concentration of NO_2 was generally low ($< 0.1 \mu\text{M}$), but a high concentration ($> 1 \mu\text{M}$) was observed in the bottom water in September, indicating the nitrification activity was high. A high concentration of NO_2 was similarly observed in the bottom water of Funka Bay, located 150 km north of Mutsu Bay, when the nitrification was active in summer (Kudo et al., 2007).

4.2 Seasonal changes in AOR

The total AOR ($V_{\text{A+B}}$) in the bottom water in Mutsu Bay was measured from spring to autumn for two years. The $V_{\text{A+B}}$ showed the maximum in September in 2018 and 2019, at which the concentrations of NO_2 and NO_3 were highest. The factors affecting ammonia oxidation reactions are reported irradiance, salinity, water temperature, pH, and dissolved oxygen concentration (Horak et al., 2013, Wang et al., 2014, He et al., 2018). Of these factors, irradiance inhibits ammonia oxidation. The water depth at the sampling site was 50 m, where the euphotic zone was 40 m, judged by 1% of the surface irradiance. Therefore, the irradiance did not influence the AOR in the bottom layer. The AOR showed a negative correlation with salinity ranging from 1 to 15 in the Yangtze estuary (Zheng et al., 2014). Salinity in the bottom layer in Mutsu Bay showed a small variation, between 33 and 34. Salinity was not the main factor affecting AOR in this study area. Ammonia oxidation activity is known to decrease with extreme pH, but there was a small seasonal variation (8.1 ± 0.1) in the seawater's pH (Mori pers. com). In addition, the dissolved oxygen concentration was high enough in the bottom waters ($\text{DO} > 2 \text{ mL L}^{-1}$).

One of the suggested effective factors for seasonal change in AOR is temperature (Kim et al., 2008). The temperature dependency of nitrification with an exponential increase in activity is found over an annual cycle observation in Narragansett Bay (Berounsky and Nixon, 1990). All known nitrifying bacteria in culture are mesophiles with optimal growth around 30 °C (Bock and Koops, 1992; Koops and Möller, 1992). Water temperatures in the bottom water during the observation period were below 13 °C from May to July, and they were between 16 and 18 °C in September. The water temperature would explain why the V_{A+B} was higher in September than in May. However, the water temperatures in the bottom water in November, when the V_{A+B} was less than 40% of that in September, were 17 °C and 16 °C in 2018 and 2019, respectively, which were not much different from those in September. This finding suggests factors besides temperature enhanced V_{A+B} and nitrification activity. We will further discuss this point in the next section.

4.2 Relationship between copy numbers and AOR, and contribution of AOA and AOB

Nakagawa et al. (2007) reported that the abundance of AOA was higher than that of AOB in the deep water of the northeast of Japan Sea. The dominance of AOA rather than AOB with one to two orders of difference in abundance was also reported in the North Sea (Wutcher et al., 2006) and off the coast of California (Santro et al., 2010). The dominance of AOA throughout the year in Mutsu Bay was consistent with these regions. The number of copies for AOA-*amoA* and AOB-*amoA* was highest in September except for AOA-*amoA* at B-0.3 m, but the number of AOA-*amoA* copies at B-0.3 m was highest in May (Fig. 9). If the number of copies for AOA-*amoA* and AOB-

1 *amoA* represents the abundance of these AOOs, the observed high ammonia oxidizing
2 activity in September seemed to be strongly influenced by the biomass of related
3 organisms.

4 To distinguish AOA and AOB activities, an incubation experiment was
5 conducted using ATU, an inhibitor of AOB. Overall, the V_A was responsible for more
6 than half of the V_{A+B} (Fig. 6), but the V_B exceeded the V_A just above the sediment (B-
7 0.3 m). The strong vertical gradient of AOR was observed in the water column within
8 several meters from the seafloor. The stratification that was formed in the bottom waters
9 in September likely maintains this vertical gradient.

10 The AOR per copy of the *amoA* gene was calculated by dividing the AOR of
11 AOA or AOB by the number of copies of AOA- or AOB-*amoA* in 24 h of culture in
12 September, when the AOR was high. To perform this calculation, the number of copies
13 per AOA cell was set to one (Walker et al., 2010; Blainey et al., 2011) and the number
14 of copies per AOB cell was set to three (Norton et al., 1996). The AOR per cell was 0.1-
15 4 fmol cell⁻¹ d⁻¹ and 0-16 fmol cell⁻¹ d⁻¹ for AOA and AOB, respectively. The AOR per
16 cell was higher for AOB than for AOA when V_B was detected. Wuchter et al. (2006)
17 reported the cellular V_A as 2-4 fmol cell⁻¹ d⁻¹. Santoro et al. (2010) reported the cellular
18 V_B as 0.2-1.5 fmol cell⁻¹ d⁻¹. The values obtained in the present study fell within the
19 range of reported values.

20 A significant correlation ($p < 0.05$) was found between the AOB-*amoA* copy
21 number at 24 h of incubation and V_B whereas no correlation was found between the
22 AOA-*amoA* copy number and V_A (Fig. 9). No correlation was found between AOR and
23 the abundance of AOA and AOB in the Arabian Sea and the central California current
24 (Santoro et al., 2010; Newell et al., 2011). However, the vertical profile of AOR was

1 correlated with the amount of AOA-*amoA* in the Gulf of California (Beman et al.,
2 2012). One of the possible reasons for these inconsistent results between the abundance
3 of AOO and AOR seemed to be that in most sequence-based studies, researchers
4 amplified DNA extracted DNA from samples, but these extracts may include DNA
5 from not only active cells but also inactive but viable cells and inactive dead cells as
6 well as extracellular DNA from lysed or degraded cells (Gaidos et al, 2011). Therefore,
7 the abundance evaluated by *amoA* copies does not always relate to the oxidation
8 activities. We conclude that not only V_A and V_B measurements but also *amoA* copy
9 number assay are necessary to better understand regeneration processes of bioavailable
10 N.

12 4.3 Influence of SS on the rate and contribution of ammonia oxidation

13 When the water at B-0.3 m was filtered with a 3- μ m filter, the total AOR
14 decreased to less than half of the AOR in the unfiltered water (Fig. 7). This decrease
15 explained that the ammonia-oxidizing microorganisms associated with SS were partially
16 removed by the filtration. Because the size of general archaea and bacteria (nitrifying
17 bacteria) is about 1-2 μ m, it is believed that free-living ammonia-oxidizing
18 microorganisms (not associated with SS) pass through a filter of 3- μ m pores. In
19 addition, the contributions of AOA and AOB to the AOR differed by filtration (Fig. 7).
20 The contribution of AOB in September 2019, when the AOR was high, accounted for
21 66% of the total AOR in the unfiltered B-0.3 m water, but it was almost zero in the
22 filtered water. However, there was no significant difference in the V_A between the
23 filtered and the unfiltered samples ($p > 0.05$). This finding suggests that AOA was free-
24 living whereas AOB was attached to the SS's surface (Fig. 10). According to Stehr et al.

(1995), there are more AOB-associated particles near the sediments than in the surface water at the mouth of the Elbe River and the abundance of AOB increases as the amount of SSs in water increases. Li et al. (2022) reported the PNR of AOA and AOB using surface sediment in the Bohai Sea and South Yellow Sea. They reported that the relative contribution of V_A ranged from 21 to 100%, but more than 50% of AOB's relative contribution was found in the summer in the Bohai Sea. These studies suggest that the surface sediment was the source of this SS. Thus, the main habitat of AOB seemed in the surface sediment and the abundance of AOB increased in summer responding to the increase in NH_4 concentration. The particle associated AOB was supplied from the surface sediment to the bottom water through resuspension and accelerated the ammonium oxidation activities in the bottom water (Fig. 10).

Conclusion

Ammonia oxidation activities increased in the bottom waters of Mutsu Bay during the summer and was accompanied by the dynamic balance reflecting the difference in the life state of AOA and AOB, free-living and particle-attached, respectively (Fig. 10). This high ammonia oxidation activity cannot be explained only by high water temperature during this period but is closely related to the increased biomass of AOBs. Ammonia oxidation by AOB accelerated the total ammonium oxidation activity in September. Ammonia oxidation occurs not only in the bottom water but also in the sediment surface layer when AOB is supplied to the bottom water in the form of suspended solids. Because ammonium oxidation is the rate-limiting step in the nutrient regeneration process in the bottom layer, including the sediment surface,

the elucidation of the ammonia oxidation process in this study is significant considering whether primary productivity can be maintained appropriately in the future. Furthermore, maintaining a healthy bottom-sediment environment is essential for sustainable bivalve aquaculture in the bay.

CRediT authorship contribution statement

Akari Aizawa: Investigation, Methodology, Writing-original draft. Yuka Watanabe: Investigation, Methodology. Kaori Hashioka: Investigation, Methodology. Aya Kadoya: Investigation, Methodology, Resources. Satoru Suzuki: Investigation, Methodology, Resources, Writing-review & editing. Takeshi Yoshimura: Investigation, Methodology, Writing-review & editing. Isao Kudo: Conceptualization, Funding acquisition, Investigation, Methodology, Supervision, Writing-review & editing.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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7 Data availability

8 Data will be available on request.

10 References

- 11 Ando, Y., Nakagawa, T., Takahashi, R., Yoshihara, K., Tokuyama, T. 2009.
12 Seasonal changes in abundance of ammonia-oxidizing archaea and ammonia-oxidizing
13 bacteria and their nitrification in sand of an eelgrass zone. *Microbes and Environments*,
14 24, 21-27. <https://doi.org/10.1264/jsme2.ME08536>
- 15 Baker, B.J., Lesniewski, R.A., Dick, G.J. 2012. Genome-enabled transcriptomics
16 reveals archaeal populations that drive nitrification in a deep-sea hydrothermal plume.
17 *The ISME journal*, 6, 2269-2279. <https://doi:10.1038/ismej.2012.64>
- 18 Beman, J.M., Popp, B.N., Alford, S.E. 2012. Quantification of ammonia
19 oxidation rates and ammonia-oxidizing archaea and bacteria at high resolution in the

Gulf of California and eastern tropical North Pacific Ocean. *Limnol. Oceanogr.*, 57,
711-726. <https://doi.org/10.4319/lo.2012.57.3.0711>

Bernhard, A.E., Tucker, J., Giblin, A.E., Stahl, D.A., 2007. Functionally distinct
communities of ammonia-oxidizing bacteria along an estuarine salinity gradient.
Environ. Microbiol. 9(6), 1439–1447. [https://doi.org/10.1111/j.1462-](https://doi.org/10.1111/j.1462-2920.2007.01260.x)
[2920.2007.01260.x](https://doi.org/10.1111/j.1462-2920.2007.01260.x).

Berounsky, V.M., Nixon, S.W. 1990. Temperature and the annual cycle of
nitrification in waters of Narragansett Bay. *Limnol. Oceanogr.*, 35, 1610-1617.
<https://doi.org/10.4319/lo.1990.35.7.1610>

Bianchi, M., Feliatra, F., Tréguer, P., Vincendeau, M.A., Morvan, J. 1997.
Nitrification rates, ammonium and nitrite distribution in upper layers of the water
column and in sediments of the Indian sector of the South Ocean. *Deep-Sea Res. II*, 44,
1017-1032. [https://doi.org/10.1016/S0967-0645\(96\)00109-9](https://doi.org/10.1016/S0967-0645(96)00109-9)

Blainey, P. C., Mosier, A. C., Potanina, A., Francis, C. A. & Quake, S. R. 2011.
Genome of a low-salinity ammonia-oxidizing archaeon determined by single-cell and
metagenomic analysis. *PLoS One* 6 (2). <https://doi.org/10.1371/journal.pone.0016626>

- 1 Bock, E., Koops, H.P. 1992. The genus *Nitrobacter* and related genera. In:
2 Balows, A., Trüper, H.G., Dworkin, M., Harder, W., Schleifer, K.H. (eds) The
3 prokaryotes, 2nd edn, Vol. III. Springer-Verlag, New York, p2302-2309.
- 4 Bouskill, M.J., Eveillard, D., Chien, D., Jayakumar, A., Ward, B.B. 2012.
5 Environmental factors determining ammonia-oxidizing organism distribution and
6 diversity in marine environments. *Environ. Microb.*, 14, 714-729.
7 <https://doi.org/10.1111/j.1462-2920.2011.02623.x>
- 8 Caffrey, J.M., Harrington, N., Solem, I., Ward, B.B. 2003. Biogeochemical
9 processes in a small California estuary. 2. Nitrification activity, community structure
10 and role in nitrogen budgets. *Mar. Ecol. Prog. Ser.*, 248, 27-40.
11 <https://doi:10.3354/meps248027>
- 12 Gaidos, E., Rusch, A., Ilardo, M., 2011. Ribosomal tag pyrosequencing of DNA
13 and RNA from benthic coral reef microbiota: community spatial structure, rare
14 members and nitrogen cycling guilds. *Environ. Microbiol.* 13 (5), 1138–1152.
15 <https://doi.org/10.1111/j.1462-2920.2010.02392.x>.
- 16 Grundle, D.S., Juniper, S.K. 2011. Nitrification from the lower euphotic zone to
17 the sub-oxic waters of a highly productive British Columbia fiord. *Mar. Chem.*, 126,
18 173-181. <https://doi.org/10.1016/j.marchem.2011.06.001>

Grasshoff, K., Kremling, K., Ehrhardt, M., 1999. Methods of Seawater Analysis.

Wiley-VCH, Weinheim, 600 pp.

Ginestet, P., Audic, J-M., Urbain, B., Block, J.-C. 1998. Estimation of nitrifying

bacterial activities by measuring oxygen uptake in the presence of the metabolic

inhibitors allylthiourea and azide. *Appl. Environm. Microbiol.*, 64, 2266-2268.

<https://doi.org/10.1128/AEM.64.6.2266-2268.1998>

Hallam, S. J., Mincer, T., Schleper, C., Preston, C., Roberts, K., Richardson, P.,

DeLong, E. 2006. Pathways of carbon assimilation and ammonia oxidation suggested

by environmental genomic analyses of marine Crenarchaeota. *PLoS Biol.* 4, 520–536.

<https://doi.org/10.1371/journal.pbio.0040437>

He, H., Zhen, Y., Mi, T.Z., Fu, L.L., Yu, Z.G., 2018. Ammonia-oxidizing

archaea and bacteria differentially contribute to ammonia oxidation in sediments from

adjacent waters of Rushan Bay, China. *Front. Microbiol.* 9, 116.

<https://doi.org/10.3389/fmicb.2018.00116>.

Horak, R., Qin, W., Schauer, A., Armbrust, E., Ingalls, A., Moffett, J., Stahl, D.,

Devol, A., 2013. Ammonia oxidation kinetics and temperature sensitivity of a natural

marine community dominated by Archaea. *ISME J.* 7, 2023–2033.

<https://doi.org/10.1038/ismej.2013.75>

Hou, J., Song, C.L., Cao, X.Y., Zhou, Y.Y., 2013. Shifts between ammonia-oxidizing bacteria and archaea in relation to nitrification potential across trophic gradients in two large Chinese lakes (Lake Taihu and Lake Chaohu). *Water Res.* 47 (7), 2285–2296. <https://doi.org/10.1016/j.watres.2013.01.042>.

Jia, Z.J., Conrad, R., 2009. Bacteria rather than archaea dominate microbial ammonia oxidation in an agricultural soil. *Environ. Microbiol.* 11, 1658–1671. <https://doi.org/10.1111/j.1462-2920.2009.01891.x>.

Kim, J.-H., Guo, X., Park, H.-S. 2008. Comparison study of the effects of temperature and free ammonia concentration on nitrification and nitrite accumulation. *Process Biochem.*, 43, 154-160. <https://doi.org/10.1016/j.procbio.2007.11.005>

Könneke, M., Bernhard, A.E., de la Torre, J.T., Walker, C.B., Waterbury, J.B., Stahl, D.A. 2005. Isolation of an autotrophic ammonia-oxidizing marine archaeon. *Nature*, 437, 543-546. <https://doi.org/10.1038/nature03911>

Koops, H.P., Möller, U.C. 1992. The lithotrophic ammonia-oxidizing bacteria. In: Balows, A., Trüper, H.G., Dworkin, M., Harder, W., Schleifer, K.H. (eds.) *The prokaryotes*, 2nd edn., Vol. III. Springer-Verlag, New York, p2625-2337.

Kudo, I., Yoshimura, T., Lee, C.W., Yanada, M., Maita, Y. 2007. Nutrient regeneration at bottom after a massive spring bloom in a subarctic coastal environment,

1 Funka Bay, Japan. J. Oceanogr., 63, 791-801. <https://doi.org/10.1007/s10872-007-0067->

2 9

3 Kudo, I., Yoshimura, M., Hashioka, K., Adachi, T., Isoda, Y. 2014. Impact of
4 bivalve culture on oligotrophication in Mutsu Bay. Bull. Coast. Oceanogr., 52, 83-92. In
5 Japanese with English abstract. https://doi.org/10.32142/engankaiyo.52.1_83

6 Li, M., He, H., Mi, T., Zhen, Y. 2022. Spatiotemporal dynamics of ammonia-
7 oxidizing archaea and bacteria contributing to nitrification in sediments from Bohai Sea
8 and South Yellow Sea, China. Sci. Tot. Environ., 825, 153972.

9 <https://doi.org/10.1016/j.scitotenv.2022.153972>

10 Nakagawa, T., Mori, K., Kato, C., Takahashi, R., Tokuyama, T. 2007.
11 Distribution of cold-adapted ammonia-oxidizing microorganisms in the deep ocean of
12 the northeastern Japan Sea. Microb. Environ., 22, 365-372.

13 <https://doi.org/10.1264/jsme2.22.365>

14 Newell, S.E., Babbitt, A.R., Jayakumar, A., Ward, B.B. 2011. Ammonia
15 oxidation rates and nitrification in the Arabian Sea. Global Biogeochem. Cycles, 25,
16 GB4016. <https://doi.org/10.1029/2010GB003940>

17 Norton, J. M., Low, J. M. & Klotz, M. G. 1996. The gene encoding ammonia
18 monooxygenase subunit A exists in three nearly identical copies in *Nitrosospira* sp.

1 NpAV. FEMS Microbiol. Lett. 139, 181–188. <https://doi.org/10.1111/j.1574->
2 6968.1996.tb08200.x

3 Olson, R.J. 1981. ^{15}N tracer studies of the primary nitrite maximum. J. Mar.
4 Res., 39, 203-226.

5 O'Mullan, G.D., Ward, B.B. 2005. Relationship of temporal and spatial
6 variabilities of ammonia-oxidizing bacteria to nitrification rates in Monterey Bay,
7 California. Appl. Environ. Microbiol., 71, 697-705.
8 <https://doi.org/10.1128/AEM.71.2.697-705.2005>

9 Painter, S.C. 2011. On the significance of nitrification with the euphotic zone of
10 the subpolar North Atlantic (Iceland basin) during summer 2007. J. Mar. Sys., 88, 332-
11 335. <https://doi.org/10.1016/j.jmarsys.2011.05.001>

12 Pedneault, E., Galand, P.E., Potvin, M., Tremblay, J.É., Lovejoy, C. 2014.
13 Archaeal amoA and ureC genes and their transcriptional activity in the Arctic Ocean.
14 Scientific reports, 4, 4661. <https://doi.org/10.1016/j.jmarsys.2011.05.001>

15 Rotthauwe, J., Witzel, K., Liesack, W. 1997. The Ammonia Monooxygenase
16 Structural Gene amoA as a Functional Marker: Molecular Fine-Scale Analysis of
17 Natural Ammonia-Oxidizing Populations. *Appl. Environ. Microbiol.*, **63**, 4704-4712.
18 <https://doi.org/10.1128/aem.63.12.4704-4712.1997>

1 Santoro, A.E., Casciotti, K.L., Francis, C.A. 2010. Activity, abundance and
 2 diversity of nitrifying archaea and bacteria in the central California Current. Environ.
 3 Microbiol., 12, 1989-2006. <https://doi.org/10.1111/j.1462-2920.2010.02205.x>
 4 Stehr, G., Böttcher, B., Dittberner, P., Rath, G. & Koops, H. P. 1995. The
 5 ammonia-oxidizing nitrifying population of the River Elbe estuary. FEMS Microbiol.
 6 Ecol. 17, 177–186. <https://doi.org/10.1111/j.1574-6941.1995.tb00141.x>
 7 Sterngren, A.E., Hallin, S. Bengtson, P. 2015. Archaeal ammonia oxidizers
 8 dominate in numbers, but bacteria drive gross nitrification in N-amended grassland soil.
 9 Front. In Microb.,6, 1350. <https://doi.org/10.3389/fmicb.2015.01350>
 10 Thamdrup, B., Fleischer, S. 1998. Temperature dependence of oxygen
 11 respiration, nitrogen mineralization, and nitrification in Arctic sediments. Aquat.
 12 Microb. Ecol., 15, 191-199. <https://doi.org/10.3354/ame015191>
 13 Walker, C., De La Torre, J., Klotz, M., Urakawa, H., Pinel, N., Arp, D.,
 14 Brochier-Armanet, C., Chain, P., Chan, P., Gollabgir, A., Hemp, J., Hügler, M., Karr,
 15 E., Könneke, M., Shin, M., Lawton, T., Lowe, T., Martens-Habbena, W., Sayavedra-
 16 Soto, L., Lang, D., Sievert, S., Rosenzweig, A., Manning, G., Stahl, D. 2010.
 17 *Nitrosopumilus maritimus* genome reveals unique mechanisms for nitrification and

1 autotrophy in globally distributed marine crenarchaea. Proc. Natl. Acad. Sci. U. S. A.
2 107, 8818–8823. <https://doi.org/10.1073/pnas.0913533107>

3 Wang, Y. F., Gu, J. D. 2014. Effects of allylthiourea, salinity, and pH on
4 ammonia/ammonium-oxidizing prokaryotes in mangrove sediment incubated in
5 laboratory microcosms. Appl. Microbiol. Biotechnol. 98, 3257–3274.
6 <https://doi.org/10.1007/s00253-013-5399-3>

7 Ward, B.B., O’Mullan, G.D. 2002. Worldwide distribution of *Nitrosococcus*
8 *oceanii*, a marine ammonia-oxidizing gamma-proteobacterium, detected by PCR and
9 sequencing of 16S rRNA and amoA genes. Appl. Environ. Microbiol., 68, 4153–4157.
10 <https://doi.org/10.1128/AEM.68.8.4153-4157.2002>

11 Wilson, K. 2001. Preparation of genomic DNA from bacteria. Current protocols
12 in molecular biology. <https://doi.org/10.1002/0471142727.mb0204s56>

13 Wutchter, C., Abbas, B., Coolen, M.J.L., Herfort, L., van Bleijswijk, J.,
14 Timmers, P., Strous, M., Teira, E., Herndl, G.J., Middelburg, J.J., Schouten, S.,
15 Sinninghe Damsté, J.S. 2006. Archaeal nitrification in the ocean. Proc. Natl. Acad. Sci.
16 USA, 103, 12317–12322. <https://doi.org/10.1073/pnas.0600756103>

17 Zheng, Y.L., Hou, L.J., Newell, S., Liu, M., Zhou, J.L., Zhao, H., et al., 2014.
18 Community dynamics and activity of ammonia-oxidizing prokaryotes in intertidal

1 sediments of the Yangtze estuary. *Appl. Environ. Microbiol.* 80 (1), 408–419.

2 <https://doi.org/10.1128/aem.03035-13>

3

Figure legends

Figure 1 Map showing sampling location (N-43) in Mutsu Bay.

Figure 2 Temperature profiles at N-43 in 2018 and 2019.

Figure 3 DIN (NH_4 , NO_2 , and NO_3) profiles at N-43 in 2018 and 2019. Shaded area designates concentration for each DIN species.

Figure 4 Spatial and temporal changes in ammonia oxidation rate (AOR) in 2018 and 2019. Error bar indicates standard error. NS = not sampled; ND = not detected.

Figure 5 Spatial and temporal changes in ammonia oxidation rate (AOR) by AOB (V_B) and AOA (V_A) in 2018 and 2019. Error bar indicates standard error. NS = not sampled; ND = not detected.

Figure 6 Relative contribution of V_A and V_B to the total AOR in September 2018 and 2019.

Figure 7 Change in the amount of SSs for unfiltered and a 3- μm filtered water (a) and change in V_B and V_A for unfiltered and a 3- μm filtered water (b). Error bar indicates standard error.

Figure 8 Spatial and temporal changes in abundances of AOA-*amoA* and AOB-*amoA* in 2018. Error bar indicates standard error. NS = not sampled.

1 Figure 9 Relationship between AOA-*amoA* abundance at 24 h and V_A (a) and
2 that between AOB-*amoA* abundance at 24 h and V_B .

3 Figure 10 Scheme explaining the role of AOA and AOB in the bottom water.
4
5

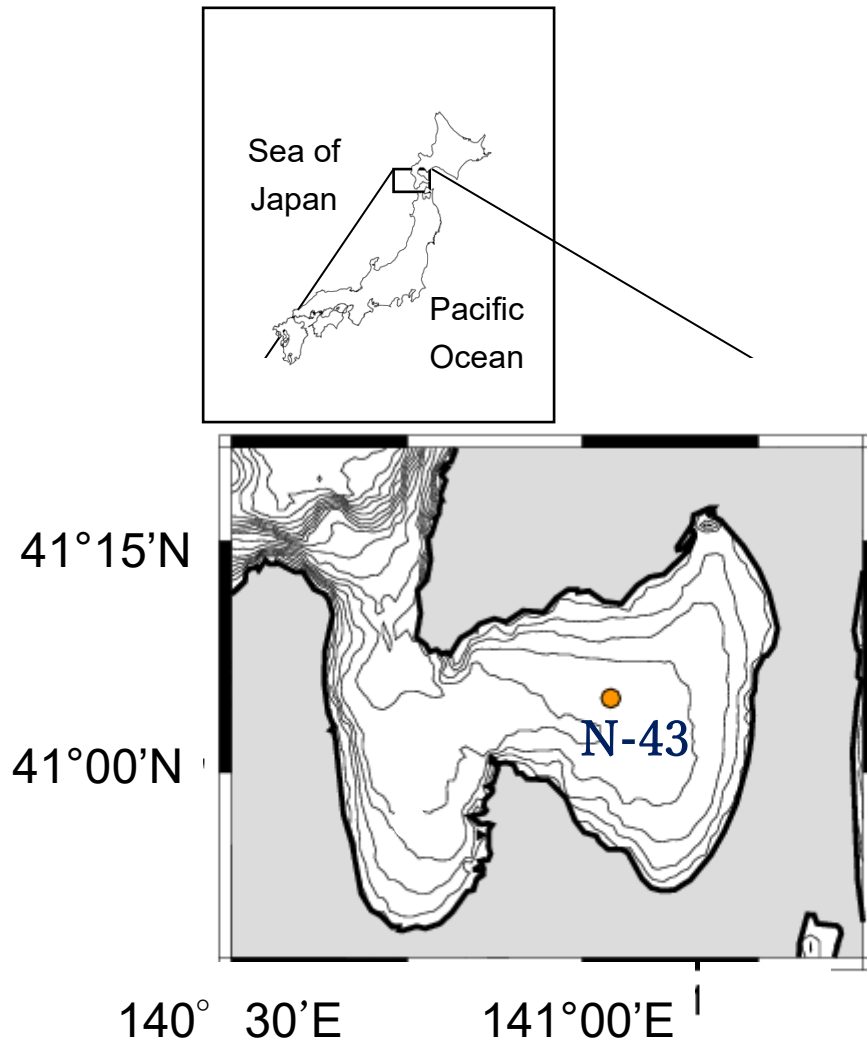


Fig. 1

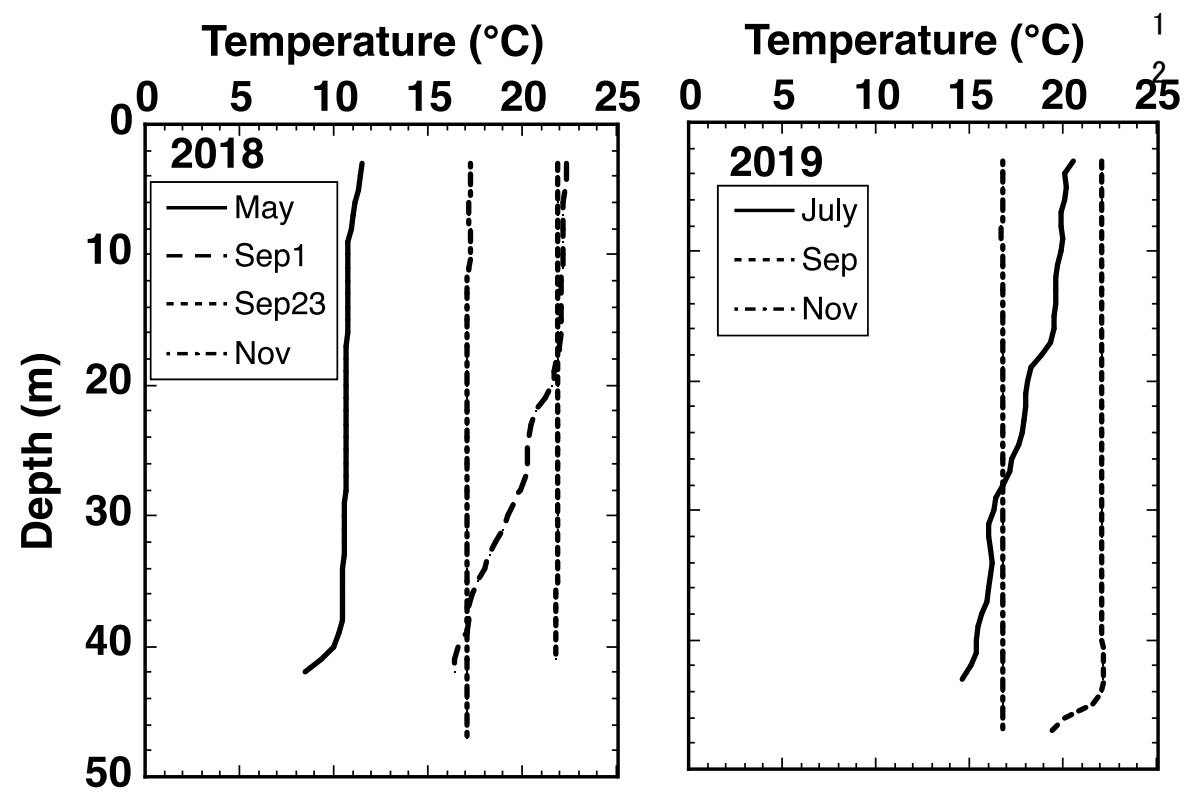


Fig. 2

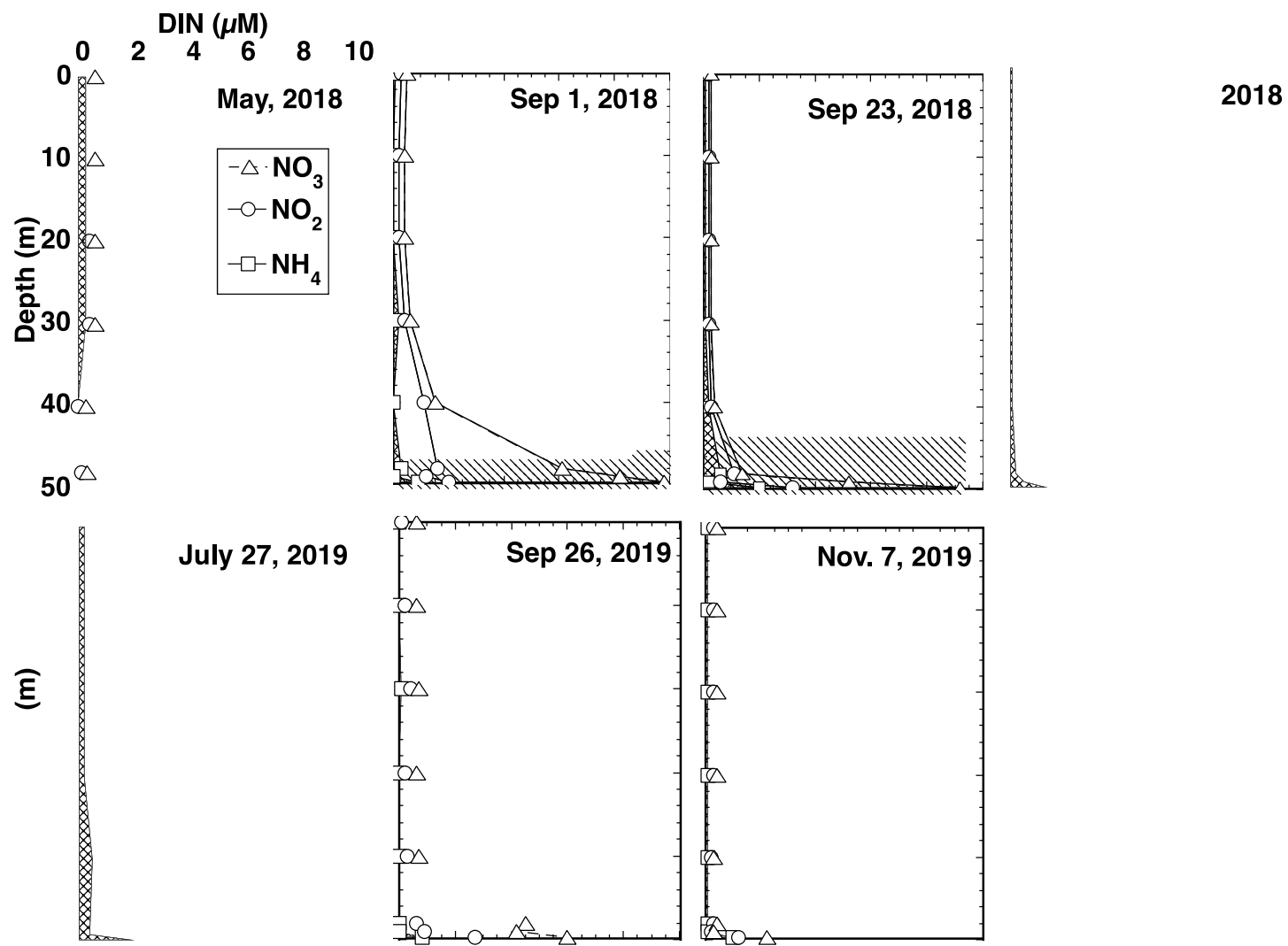


Fig. 3

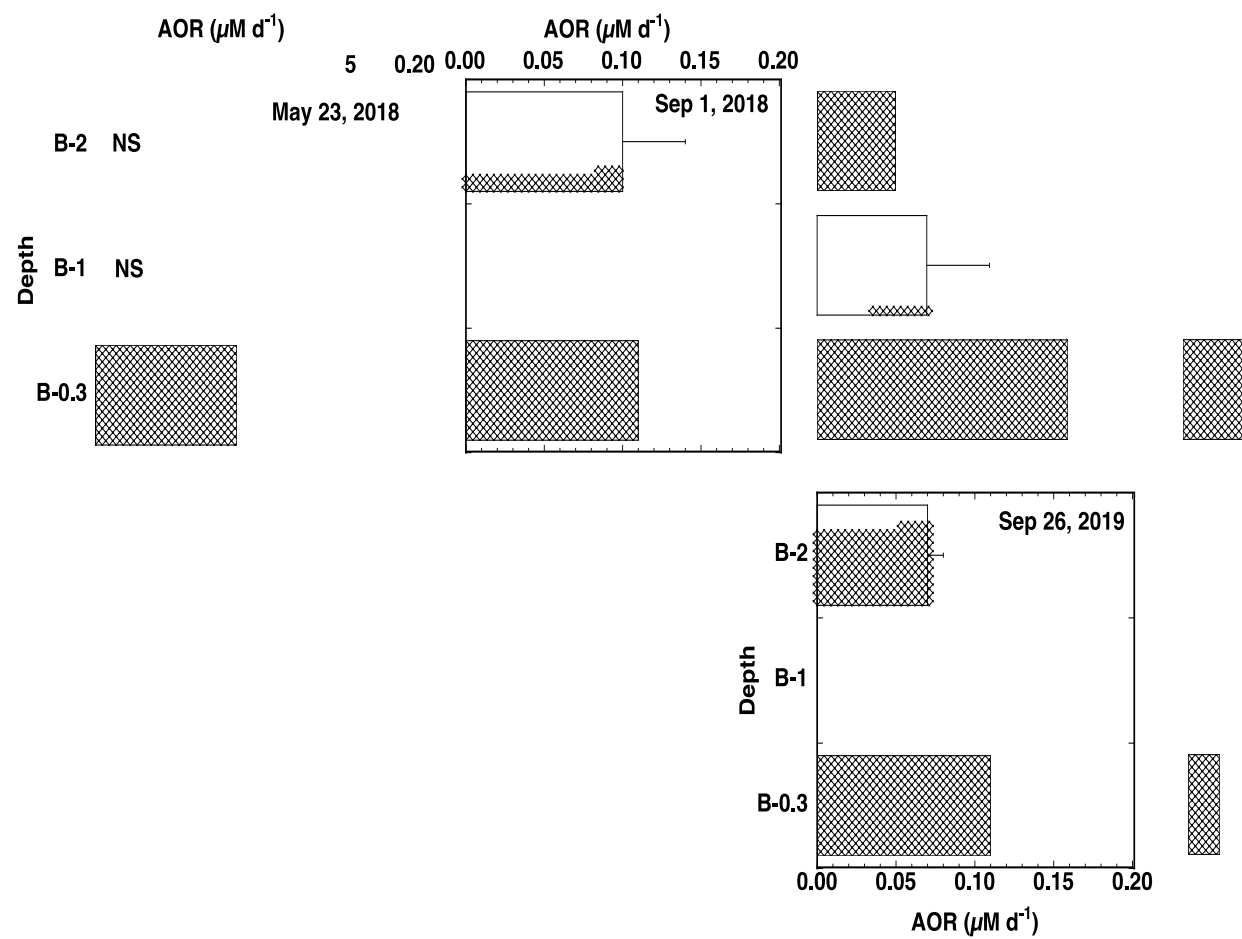


Fig. 4

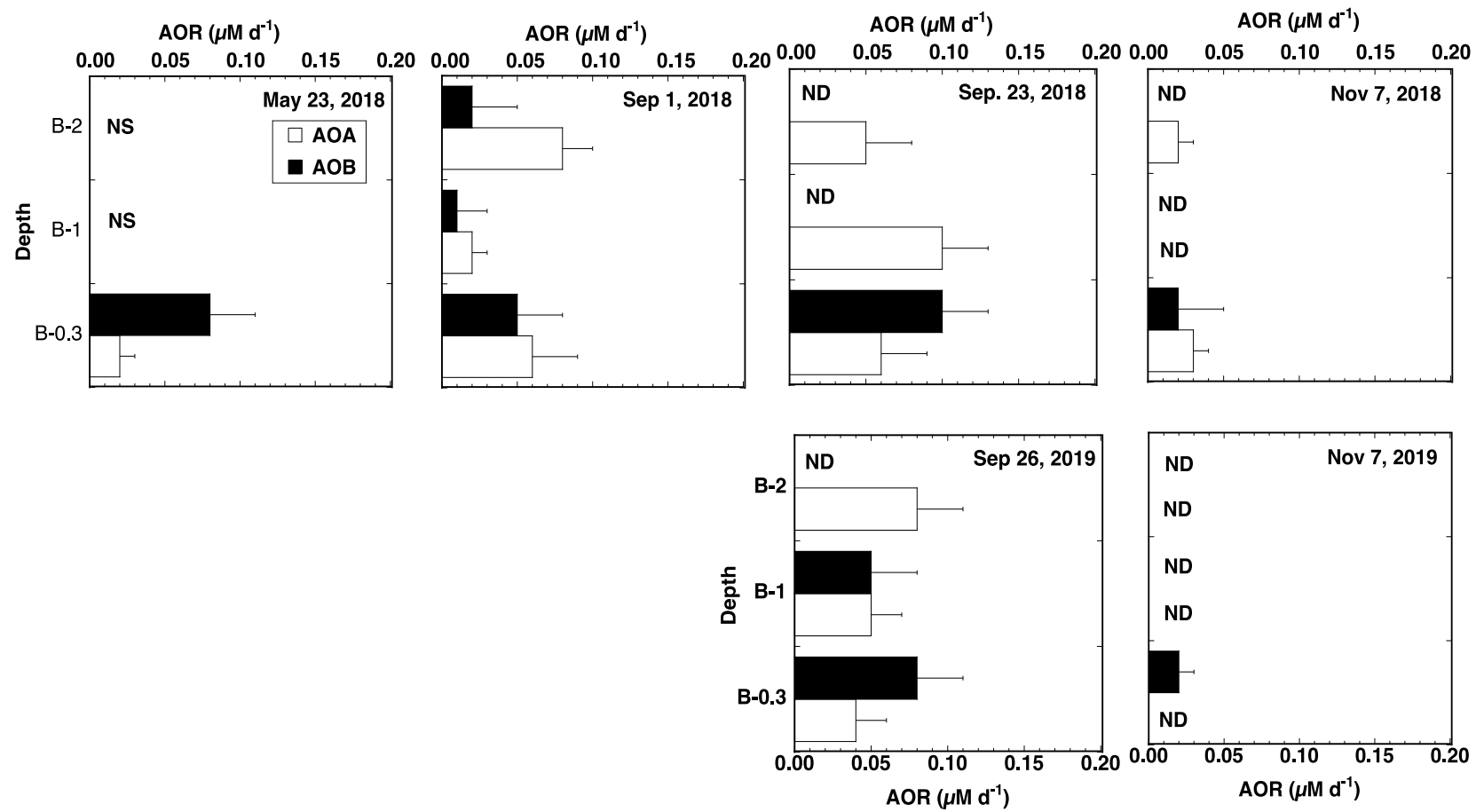


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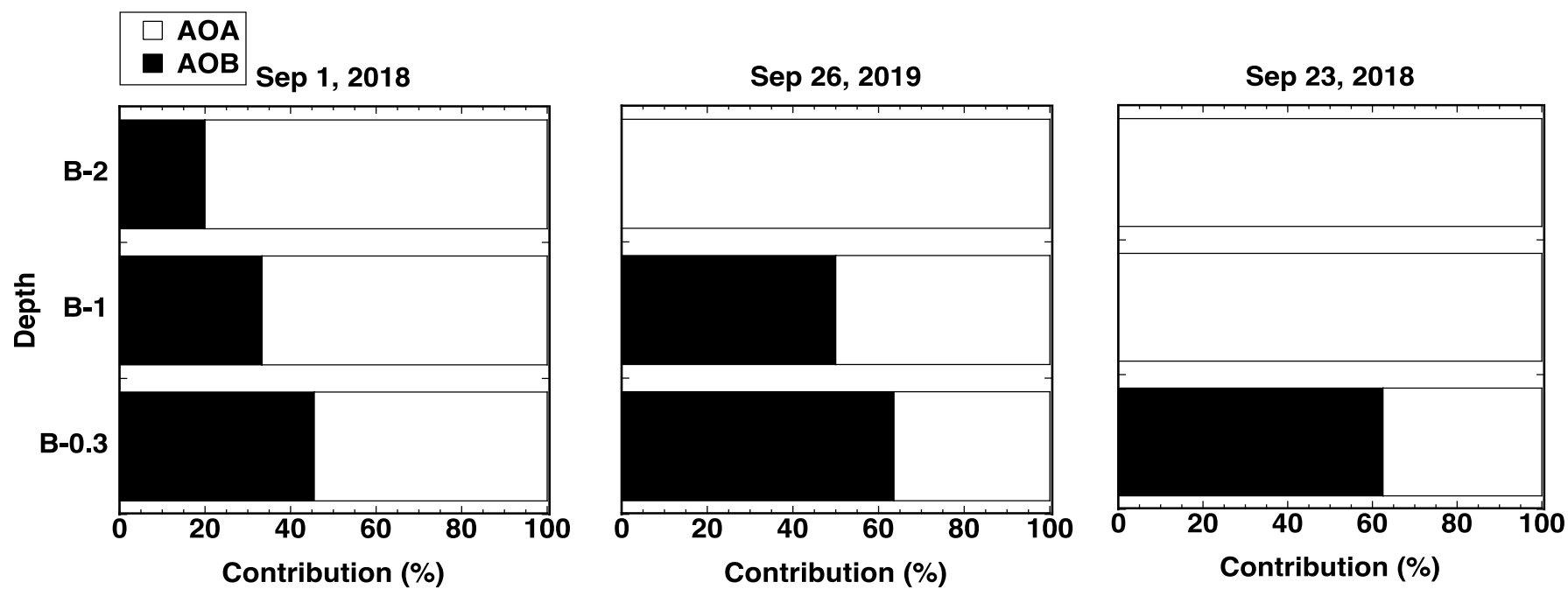


Fig. 6

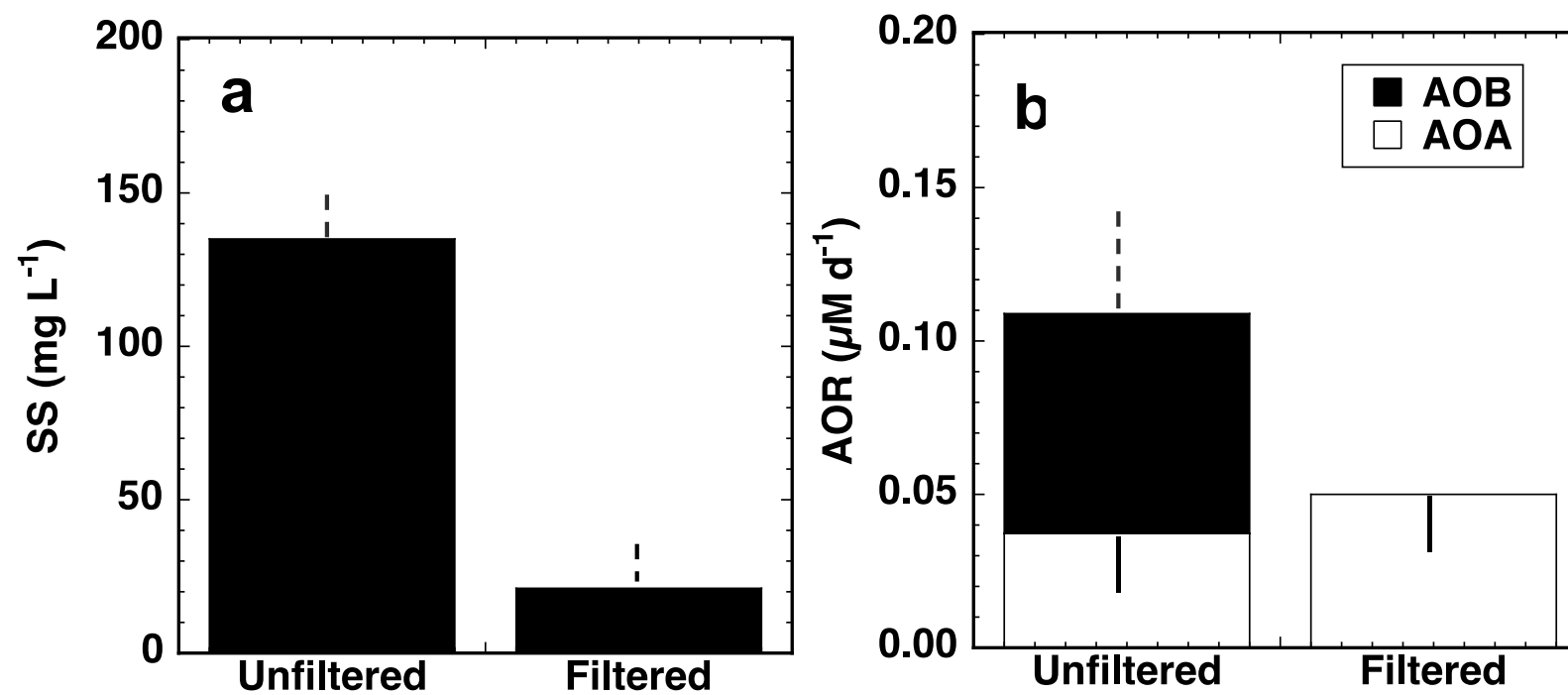


Fig. 7

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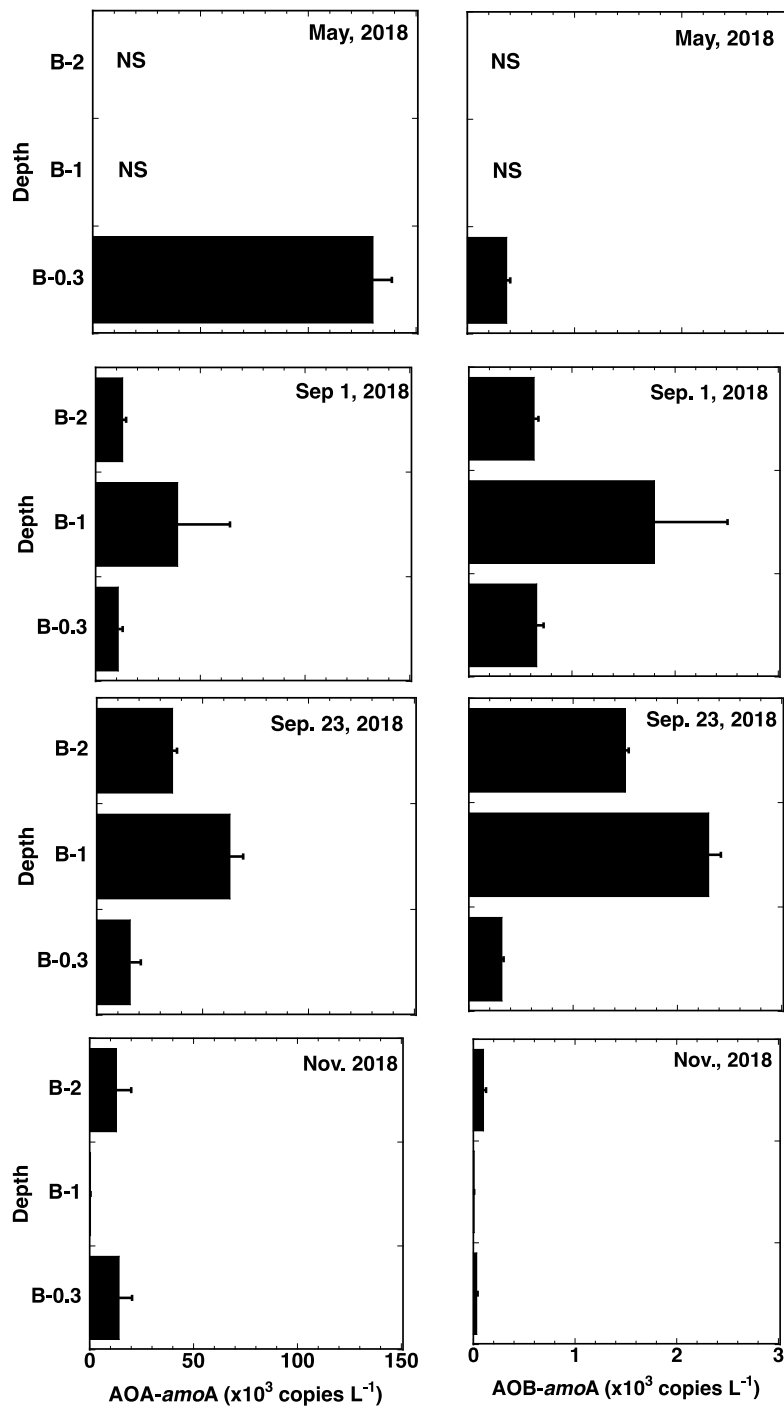


Fig. 8

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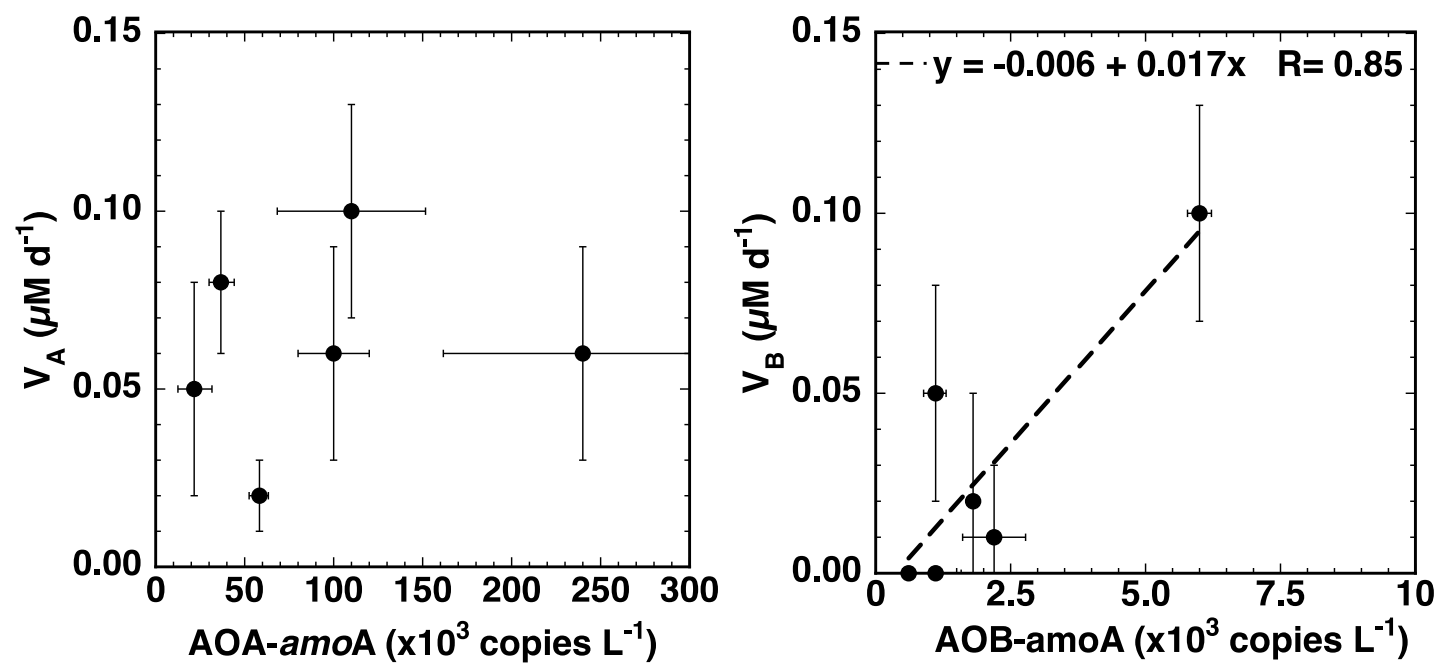


Fig. 9

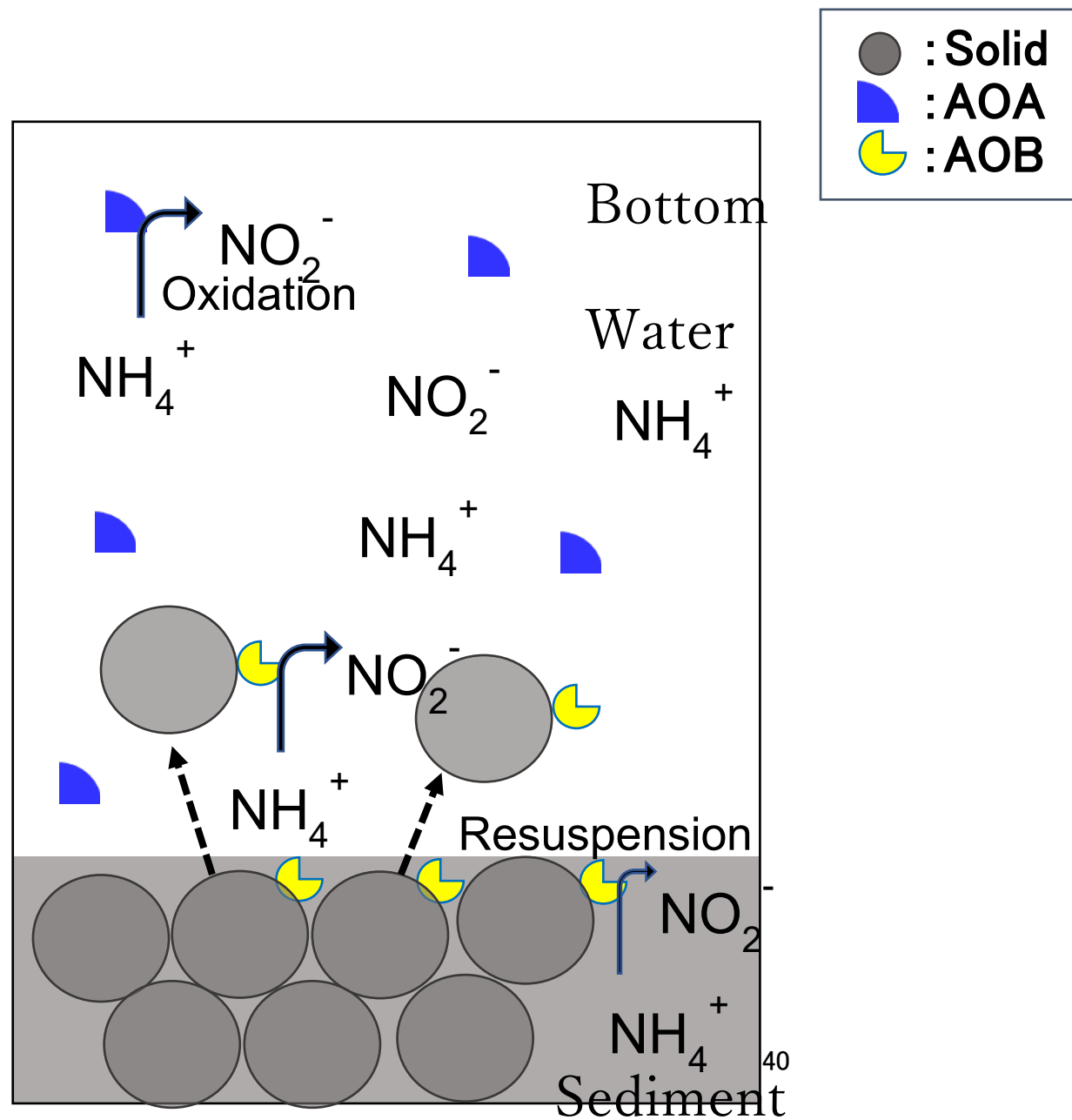


Fig. 10

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Table 1 List of frequently used acronyms

AOA	Ammonia Oxidizing Archaea
AOB	Ammonia Oxidizing Bacteria
AOOs	Ammonia Oxidizing Organisms
AOR	Ammonia Oxidation Rate
ATU	Allyl-thiourea
DIN	Dissolved Inorganic Nitrogen
PCR	Polymerase Chain Reaction
PNR	Potential Nitrification Rate
SS	Suspended Solids

2

Table 2 Primers for q-PCR analysis

Gene	Primer Set	Sequence 5'-3'	Length	Reference
AOA-amoA	crenAMO_F	ATGGTCTGGCTAAGACGMTGTA	646 bp	Hallam et al. (2006)
	crenAMO_R	CCCACTTTGACCAAGCGGCCAT		
AOB-amoA	AmoA-1F	GGGGTTTCTACTGGT	491 bp	Rotthauwe et al. (1997)
	AmoA-2R	CCCCTCKGSAAAGCCTTCTTC		