

**Intact polar lipids of Thaumarchaeota and anammox  
bacteria as indicators of N-cycling in the Eastern Tropical  
North Pacific oxygen deficient zone**

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1 **Abstract**

2 In the last decade our understanding of the marine nitrogen cycle has improved considerably  
3 thanks to the discovery of two novel groups of microorganisms: ammonia-oxidizing archaea  
4 (AOA) and anaerobic ammonia-oxidizing (anammox) bacteria. Both groups are important in  
5 oxygen deficient zones (ODZs), where they substantially affect the marine N-budget. These  
6 two groups of microbes are also well known for producing specific membrane lipids, which  
7 can be used as biomarkers to trace their presence in the environment. We investigated the  
8 occurrence and distribution of AOA and anammox bacteria in the water column of the Eastern  
9 Tropical North Pacific (ETNP) ODZ, one of the most prominent ODZs worldwide.  
10 Suspended particulate matter (SPM) was collected at different depths of the water column in  
11 high resolution, at both a coastal and an open ocean setting. The SPM was analyzed for AOA-  
12 and anammox bacteria-specific intact polar lipids (IPLs), i.e. hexose-phosphohexose (HPH)-  
13 crenarchaeol and phosphatidylcholine (PC)-monoether ladderane. Comparison with oxygen  
14 profiles reveals that both the microbial groups are able to thrive at low (<1  $\mu\text{M}$ )  
15 concentrations of oxygen. Our results indicate a clear niche segregation of AOA and  
16 anammox bacteria in the coastal waters of the ETNP, but a partial overlap of the two niches of  
17 these microbial species in the open water setting. The latter distribution suggests the potential  
18 for an interaction between the two microbial groups at the open ocean site, although the  
19 nature of this hypothetical interaction (i.e. either competition or cooperation) remains unclear.  
20

## 1 **1. Introduction**

2 The marine nitrogen cycle has been widely investigated, as nitrogen is one of the main  
3 limiting factors of primary production in the upper sunlit layers of the oceans (Arrigo, 2005;  
4 Codispoti L.A., 1997) and the ocean accounts for about half of the global net primary  
5 production (Field et al., 1998; Gruber and Galloway, 2008). In the traditional view, the marine  
6 nitrogen cycle includes nitrogen fixation as the main input of nitrogen in the ocean and  
7 dinitrogen gas formed by denitrification as the main output, so that these two pathways are  
8 mainly responsible for the marine nitrogen budget status (Karl et al., 1997). In 2001,  
9 Codispoti and coauthors suggested that in the present-day ocean, the nitrogen budget is not in  
10 a steady state, but rather out of balance, with denitrification fluxes being underestimated  
11 (Codispoti et al., 2001). Nitrogen fixation is mediated by few microorganisms, including  
12 cyanobacteria, while denitrification is performed by a wide range of microorganisms with  
13 different metabolic features, able to switch from aerobic to anaerobic nitrate ( $\text{NO}_3^-$ ) dependent  
14 respiration modes (Lam and Kuypers, 2011). In this classical view, nitrification, representing  
15 the major oxidative part of the cycle, connecting organic nitrogen to  $\text{NO}_3^-$  (Codispoti et al.,  
16 2001; Lam and Kuypers, 2011), was seen exclusively as an aerobic process carried out by  
17 ammonia- (AOB), and nitrite-oxidizing bacteria, members of the  $\beta$ - and  $\gamma$ -proteobacteria. The  
18 nitrification reaction is divided into two steps, performed by distinct bacterial groups. In the  
19 first part ammonium ( $\text{NH}_4^+$ ) is oxidized to nitrite ( $\text{NO}_2^-$ ), whereas in the second  $\text{NO}_2^-$  is  
20 oxidized to nitrate ( $\text{NO}_3^-$ ). In both cases oxygen serves as electron acceptor, although AOB  
21 have been reported to perform nitrification in sub-oxic conditions (Lam et al., 2007; Schmidt  
22 and Bock, 1997).

23 The overall understanding of the marine nitrogen cycle has substantially changed in  
24 the last decade (Fig. 1). Specific archaea were discovered to be important players in the  
25 marine nitrogen cycle (Venter et al., 2004) as some of them perform nitrification in the marine

1 water column and sediment (Francis et al., 2005; Könneke et al., 2005; Wuchter et al., 2006).  
2 The group of archaea capable of nitrification has recently been relocated in a separate phylum  
3 named Thaumarchaeota (Brochier-Armanet et al., 2008; Spang et al., 2010). Compared to  
4 their bacterial counterpart, ammonia oxidizing archaea (AOA) are often more abundant in the  
5 ocean (Karner et al., 2001; Lam et al., 2007; Wuchter et al., 2006), accounting for 20% of  
6 picoplankton and 40% of the estimated total number of cells (Karner et al., 2001). These  
7 microorganisms are able to cope with low oxygen conditions (Coolen et al., 2007; Lam et al.,  
8 2007; Park et al., 2010; Pitcher et al., 2011b; Sinninghe Damsté et al., 2002a), have low  
9 substrate requirements (Martens-Habbena et al., 2009) and are able to utilize a highly energy  
10 efficient CO<sub>2</sub>- fixation pathway (Könneke et al., 2014); new coastal marine AOA isolates  
11 show obligate mixotrophy and vary in their adaptive ability to different environmental  
12 parameters (Qin et al., 2014). All these features have been suggested to provide a reason for  
13 AOA observed dominance over ammonia oxidizing bacteria (AOB) as ammonia oxidizers in  
14 the open oceans (Könneke et al., 2014; Pester et al., 2011). Moreover, a 'novel' process in the  
15 nitrogen cycle, named anammox, was discovered. Anaerobic ammonia oxidizing (anammox)  
16 bacteria, are a unique group of microorganisms member of the order of *Planctomycetales*  
17 (Strous et al., 1999). They are able to oxidize ammonium (NH<sub>4</sub><sup>+</sup>) to molecular nitrogen (N<sub>2</sub>)  
18 under anoxic conditions, using nitrite (NO<sub>2</sub><sup>-</sup>) as electron acceptor (van de Graaf et al., 1995).  
19 Anammox bacterial activity has been detected in marine anoxic sediments and waters  
20 (Dalsgaard et al., 2003; Kuypers et al., 2003; Thamdrup and Dalsgaard, 2002) and has been  
21 recognized to contribute, along with denitrifying bacteria, to the loss of N<sub>2</sub> from the ocean  
22 (Galán et al., 2009; Hamersley et al., 2007; Kuypers et al., 2005; Lam et al., 2009; Thamdrup  
23 et al., 2006). Despite different oxygen tolerances, anammox bacteria and Thaumarchaeota  
24 have been observed to co-exist in different settings, particularly in oxygen deficient zones  
25 (ODZs) and anoxic waters (Coolen et al., 2007; Francis et al., 2005; Lam et al., 2007; Pitcher

1 et al., 2011b; Woebken et al., 2007). These two microbial groups can potentially benefit from  
2 each other, because the thaumarchaeotal nitrification might be coupled with the anammox  
3 process by providing the  $\text{NO}_2^-$  anammox bacteria need and, at the same time, consume  
4 oxygen to which anammox bacteria are sensitive. Alternatively, when nitrite is provided to  
5 anammox by other sources, the two groups might compete for  $\text{NH}_4^+$  (Yan et al., 2012).

6 In this study we investigated the occurrence and depth distribution of Thaumarchaeota  
7 and anammox bacteria in the Eastern Tropical North Pacific (ETNP) oxygen deficient zone,  
8 one of the most extended ODZs in the contemporary ocean. The presence of AOA and  
9 anammox bacteria has been reported in the ETNP oxygen deficient zone by a few studies  
10 (Beman et al., 2008, 2012, 2013; Francis et al., 2005; Podlaska et al., 2012; Rush et al., 2012)  
11 and the significance of the two microbial groups to local marine nitrogen cycling is starting to  
12 be elucidated for other ODZs (Dalsgaard et al., 2003; Galán et al., 2009; Kalvelage et al.,  
13 2013; Kuypers et al., 2003, 2005; Lam et al., 2007, 2009; Pitcher et al., 2011b; Ward et al.,  
14 2009). However, the spatial distribution and the possible co-occurrence of the two groups in  
15 the ETNP have not been investigated in detail as well as the relative contribution of AOA and  
16 anammox to the local N-cycle and their possible interactions. To the best of our knowledge  
17 only one study so far has examined concurrently the presence of the two microbial groups in  
18 the southern part of the ETNP (Podlaska et al., 2012). Other studies on the ETNP ODZ have  
19 investigated the presence of AOA along a north-south transects following the coastal line of  
20 southern California (Beman et al., 2008, 2012, 2013) and the occurrence of anammox bacteria  
21 in the southern ETNP ODZ (Rush et al., 2012). These studies did not investigate the  
22 occurrence of AOA and anammox bacteria at true open ocean sites and a comparison of AOA  
23 and anammox bacteria dynamics between coastal and open ocean waters is still missing. To  
24 fill this gap in the current knowledge we performed high resolution water sampling, at both  
25 coastal and open ocean settings in the ETNP ODZ.

1           To trace the two microbial groups we applied intact polar lipids (IPLs) specific for  
2 these groups, which have proved to be good biomarkers in various settings (Bale et al., 2013;  
3 Buckles et al., 2013; Lengger et al., 2012; Pitcher et al., 2011b, 2011c). Anammox bacteria  
4 produce unique ladderane fatty acids which contain 3-5 concatenated cyclobutane moieties  
5 (Sinninghe Damsté et al., 2002c). They are attached to the glycerol backbone with polar head  
6 groups comprising phosphocholine (PC) and phosphoethanolamine (PE) (Boumann et al.,  
7 2006; Rattray et al., 2008). Thaumarchaeota produce also specific biomarker lipids, i.e.  
8 crenarchaeol, a glycerol dibiphytanyl glycerol tetraether (GDGT) lipid, containing a  
9 cyclohexane moiety beside four cyclopentane moieties (de la Torre et al., 2008; Pitcher et al.,  
10 2010; Schouten et al., 2008; Sinninghe Damsté et al., 2002b, 2002c). Attached to crenarchaeol  
11 are various polar headgroups such as monohexose (MH), dihexose (DH) and a hexose-  
12 phosphohexose (HPH) (Schouten et al., 2008) with the latter being the most suitable for  
13 tracing living active cells (Pitcher et al., 2011a). By applying these specific IPLs, i.e. HPH-  
14 crenarchaeol and PC-monoether ladderane, respectively, we investigate the depth habitat of  
15 the Thaumarchaeota and anammox bacteria in the ETNP oxygen deficient zone and the  
16 factors controlling their ecological niche.

17

## 1 **2. Materials and methods**

### 2 **2.1. Environmental setting of the ETNP**

3 The ETNP ODZ is one of the thickest in the contemporary ocean and extends to depths as  
4 deep as ~1000 m. Geographically it ranges from ~25°N (i.e. Baja California) to ~10°N (i.e.  
5 Costa Rica) and from ~160°W in the North Pacific Ocean to the coast of Mexico and Costa  
6 Rica. It is a permanent feature of the eastern tropical Pacific region (Paulmier and Ruiz-Pino,  
7 2009). The region is important for its role in the global carbon cycle, for its involvement in El  
8 Niño Southern Oscillation, and it is economically relevant for fishery (Fiedler and Lavín,  
9 2006). A shallow and strong thermocline causes water stratification and weak exchanges of  
10 nutrients and oxygen between surface waters and sub-thermocline layers, which are poorly  
11 ventilated (Lavín et al., 2006). This feature is further exacerbated by Ekman pumping which  
12 causes coastal and open ocean upwelling (Lavín et al., 2006). The hydrology of the eastern  
13 tropical Pacific is influenced by water circulation features and by strong winds in the part  
14 close to the American continent (Kessler, 2006). The ETNP oxygen deficient zone comprises  
15 part of the North Pacific sub-tropical gyre, specifically it is delimited southeastern by the  
16 California Current (CC), the North Equatorial Current (NEC) and the North Equatorial  
17 Countercurrent (NECC) (Karstensen et al., 2008). The boundary area where the CC, flowing  
18 along the coast of the Baja California and southern, encounters the NEC is characterized by  
19 Ekman transport westward and upwelling mainly off the Californian coast and in a weakened  
20 magnitude off the north Mexico coast (Kessler, 2006; Lavín et al., 2006). Our sampling area  
21 lies in this transition region; however the upper water circulation in this region is not fully  
22 understood yet (Kessler, 2006). Although ODZs have been now studied for almost a century,  
23 only recently it has been possible to determine in situ concentration in these areas to the  
24 accuracy of nanomolar O<sub>2</sub> (Revsbech et al., 2009). This has allowed to prove that ODZ  
25 regions, including the ETNP, are functionally devoid of oxygen, although respiratory rates

1 indicate that aerobic metabolisms successfully occur, even in such extreme conditions  
2 (Canfield et al., 2010; Jensen et al., 2011; Revsbech et al., 2009; Thamdrup et al., 2012; Tiano  
3 et al., 2014).

4

## 5 **2.2. Sampling**

6 Sampling was performed at twelve stations during the Eastern Tropical North Pacific  
7 (TN278) cruise (R/V Thomas G. Thompson, March-April 2012) (Fig. 2. panel A). The cruise  
8 route was split in two legs with the former one comprising six sampling stations (Fig 2. panel  
9 B), in very close proximity to each other, in coastal waters, north-west of the departure port of  
10 Manzanillo (Colima, MX), and the latter including six sampling stations, clustered closely  
11 together, in open ocean waters south-west of the departure port, around the area known as the  
12 Moctezuma Trough (Fig. 2. panel C). Suspended particulate matter (SPM) samples were  
13 collected on pre-ashed 0.7  $\mu\text{m}$  pore size glass fiber (GF) filters, mounted in McLane WTS-LV  
14 in situ filtration systems. At each sampling site four McLane pumps were deployed  
15 simultaneously at different depths (Table 1). The volume of water filtered varied according to  
16 the depth and the material collected (Table 1). Upon the recovery of the pumps the GF filters  
17 were removed, split in two halves and frozen at  $-40^{\circ}\text{C}$ .

18 Physical parameters of the water column were recorded by conductivity-temperature-  
19 density (CTD) equipment (SBE-911, Sea-Bird Electronics); dissolved-oxygen depth  
20 concentrations were measured by a SBE 43 electrochemical sensor mounted on the CTD  
21 rosette. Sensor oxygen concentrations were calibrated against on-deck Winkler titrations. The  
22 data reported here do not take into account recent evidence that these techniques (i.e. Clark  
23 electrodes, Winkler titrations) overestimate oxygen at the very lowest concentrations (Tiano  
24 et al., 2014). Water samples for inorganic nutrient profiles were collected using 24 $\times$ 10 L  
25 Niskin bottles mounted on a rosette to the CTD. The CTD was cast shortly before or after the



1 deployment of the McLane pumps. In case this was not possible, data from another station,  
2 the closest in time and space to that of the deployed in situ pumps, were used. This means that  
3 at some sites the depths sampled for nutrients data do not always directly correspond to the  
4 depths at which SPM was sampled with the in situ pumps (Table 1). The detection limits for  
5  $\text{NO}_3^-$ ,  $\text{NO}_2^-$ , and  $\text{NH}_4^+$  were respectively 0.08  $\mu\text{M}$ , 0.01  $\mu\text{M}$  and 0.07  $\mu\text{M}$ . The  
6 electrochemical oxygen sensor SBE 43 has a detection limit of 1-2  $\mu\text{M}$  (Tiano et al., 2014).

7

### 8 **2.3. Intact polar lipid analysis**

9 Intact polar lipids were extracted from freeze-dried SPM filter halves using a modified Bligh-  
10 Dyer technique as described in Sturt et al. (2004) with some adjustments as described in  
11 Schouten et al. (2008). Briefly, a known volume of methanol (MeOH):dichloromethane  
12 (DCM):phosphate buffer (P-buffer) (2:1:0.8, v/v/v) was added to the filter in a glass  
13 centrifuge tube and the total lipid contents were extracted in a sonication bath for 10 min.  
14 After centrifugation for 3 min at 2000 rpm the supernatant was removed. The extraction was  
15 repeated two more times and the supernatants combined. To induce separation of the  
16 combined supernatant into two phases, additional DCM and P-buffer were added to a new  
17 volume ratio of 1:1:0.9 DCM:MeOH:P-buffer. The mixture was centrifuged for 2 min at 3000  
18 rpm after which the DCM layer was removed. The procedure was repeated two more times  
19 and the combined DCM phases were collected in a round bottom flask, reduced under rotary  
20 vacuum and completely dried under  $\text{N}_2$  (Schouten et al., 2008; Sturt et al., 2004).

21 IPLs were analyzed directly in the extract using a high performance liquid  
22 chromatography (HPLC)-electrospray ionization (ESI)/triple quadrupole MS in selected  
23 reaction monitoring (SRM) mode as described by Pitcher et al. (2010). In order to minimize  
24 possible variations in the IPLs response factors, the extract was analyzed in the same batch.  
25 Briefly, an Agilent (Palo-Alto, CA, US) 1100 series LC equipped with a thermostat-controlled

1 auto-injector was used coupled to a Thermo TSQ Quantum EM triple quadrupole MS  
2 equipped with an Ion Max source with ESI probe. The SRM method for the crenarchaeol IPLs  
3 was targeting specifically HPH-crenarchaeol (Schouten et al., 2008; Pitcher et al., 2010). Due  
4 to the lack of a standard, HPH-crenarchaeol was quantified as the integrated IPL area peak  
5 response units  $L^{-1}$  (i.e. r.u.  $L^{-1}$ ), unveiling the relative depth distribution of the lipid biomarker  
6 in the water column, but not providing information on its absolute abundance. The anammox-  
7 specific membrane lipid  $C_{20}$ -[3]-ladderane with a phosphocholine (PC)-monoether was  
8 analyzed according to Jaeschke et al. (2009). The intact ladderane monoether lipid was  
9 quantified referring to an external calibration curve of an isolated  $C_{20}$ -[3]-ladderane (PC)-  
10 monoether standard (Jaeschke et al., 2009).

## 1 **3. Results**

### 2 **3.1. Oxygen and nutrient profiles**

3 During the ETNP (TN278) cruise in March-April 2012 the water column of the ETNP ODZ  
4 was sampled at high resolution for SPM, at depths from 20 to 2000 m in coastal waters and  
5 from 50 to 2500 m in open ocean waters at a number of geographically nearby stations (Fig.  
6 2; Table 1). To compile all our data in two (coastal and open ocean) composite profiles, we  
7 report our nutrient (Figs. 3a-c; 3g-i), oxygen (Figs. 3d and 3l), and lipid SPM (Figs. 3e and 3f;  
8 3m and 3n) data relative to the potential density anomaly,  $\sigma_\theta$  ( $\text{kg m}^{-3}$ ) of the water masses  
9 sampled at each of the coastal or open ocean stations, respectively. In the upper water column  
10 values for  $\sigma_\theta$  differ at the two sampling sites as a consequence of differences in salinity (data  
11 not shown).

12 For both locations, the oxygen profiles (Figs. 3d and 3l) obtained by an oxygen  
13 electrochemical sensor mounted on the CTD of the nearby stations were virtually identical.  
14 Station 106, which is slightly further away from the other coastal stations (Fig. 2), is an  
15 exception since the upper oxycline is located at a  $\sigma_\theta$  of 25.5, ca. 30 m deeper than at the other  
16 stations. For the other coastal stations the oxycline occurs in shallow waters at a  $\sigma_\theta$  of 25.0,  
17 i.e. ~30 m depth, whilst for the in open ocean waters it is located at  $\sigma_\theta$  25.5, which  
18 corresponds to ~100 m depth. Because of the virtual identical oxygen-  $\sigma_\theta$  profiles, it was  
19 decided that all the data for the coastal (except for station 106) and open ocean stations could  
20 be combined in presenting the nutrient concentrations and IPL data from SPM. In this way we  
21 provide an expanded view of the vertical distribution of the two microbial species along the  
22 water column.

23 In the upper coastal waters the oxygen concentration varies between 250 to 50  $\mu\text{M}$   
24 until  $\sigma_\theta$  23.5 (i.e. ~15-20 m), then drops to values below 20  $\mu\text{M}$  at the upper oxycline (Fig.  
25 3d). In this setting, the ODZ spans from  $\sigma_\theta$  25.0 to 27.0 (i.e. between ~35 to ~800 m), with a

1 minimal oxygen concentration below 1  $\mu\text{M}$  in the core (Fig. 3d) (see also Tiano et al., 2014).  
2 Below 850 m  $\sigma_\theta$  is between 27.0 and 27.5 and the oxygen concentration increases gradually  
3 again until  $\sim 75 \mu\text{M}$ , at 2000 m depth (Fig. 3d). In the open ocean waters oxygen  
4 concentration is stable at  $\sim 200 \mu\text{M}$  until  $\sigma_\theta$  22.5, at  $\sim 55$  m depth, and rapidly decreases to  
5 values close to 1  $\mu\text{M}$  at  $\sigma_\theta$  25.5, at 100 m depth. The ODZ extends until  $\sigma_\theta$  27.0, at 850 m  
6 depth. Below the lower oxycline the oxygen concentration increases again to  $\sim 120 \mu\text{M}$  at  
7 3000 m depth, where  $\sigma_\theta$  ranged between 27.5 and 28.0 (Fig. 3l).

8 Nutrient concentration data (i.e.  $\text{NO}_3^-$ ,  $\text{NO}_2^-$ ,  $\text{NH}_4^+$ ) from different stations were  
9 combined in one coastal and one open ocean setting, as described above for the oxygen depth  
10 profiles. The resulting profiles show distinct patterns (Figs. 3a-c; 3g-i). In both coastal and  
11 open waters nitrate is the most abundant nitrogen species and show two maxima at different  
12  $\sigma_\theta$  (Figs. 3a and 3g). In coastal waters the first maximum of  $\sim 25 \mu\text{M}$  occurs at the upper  
13 oxycline at  $\sigma_\theta$  25, then the concentrations decrease to 15-20  $\mu\text{M}$  until  $\sigma_\theta$  26.5. The second  
14 maximum of  $\sim 45 \mu\text{M}$  occurs at  $\sigma_\theta$  27.4, just below the lower oxycline (i.e. at 1000 m depth)  
15 (Fig. 3a). In open ocean waters the trend in nitrate appears similar to the one in the coastal  
16 waters, although some differences can be noticed (Fig. 3g). The first maximum (i.e.  $\sim 25 \mu\text{M}$ )  
17 in this case is broader and deeper, spanning from  $\sigma_\theta$  24.5 to 26, where the upper oxycline is  
18 located. The second deeper maximum (i.e.  $\sim 50 \mu\text{M}$ ) as well occurs where the waters start to  
19 be re-oxygenated (i.e.  $\sigma_\theta$  27.4, depth 1100 m). Like for nitrate, the profiles of nitrite are  
20 slightly different for the coast and open ocean waters (Figs. 3b and 3h). In the former setting  
21 the shallow maximum occurs at  $\sigma_\theta$  24.6, at declining oxygen concentrations. The peak is only  
22  $\sim 1 \mu\text{M}$  and rather narrow. The lower peak of  $\sim 8 \mu\text{M}$  is located at  $\sigma_\theta$  26 (i.e. 88 m depth) in the  
23 core ODZ (Fig. 3b). The first nitrite maximum in the open ocean reaches 1 to 2  $\mu\text{M}$  and spans  
24 from fully oxygenated waters to the oxycline (i.e.  $\sigma_\theta$  from  $\sim 23$  to  $\sim 25$ ). The deeper maximum  
25 on the other hand (i.e.  $\sim 6 \mu\text{M}$ ) occurs in the upper ODZ (i.e.  $\sigma_\theta$  26.2) (Fig. 3h). Finally, at

1 both sampling sites  $\text{NH}_4^+$  concentrations are mostly below the detection limit of  $0.07 \mu\text{M}$   
2 (Figs. 3c and 3i) with the exception of a few data points at the costal oxycline (i.e.  $\sigma_\theta$  24.6)  
3 and in the open water where the oxygen decreases, and the  $\text{NH}_4^+$  concentrations are  
4 respectively  $0.6 \mu\text{M}$  and  $\sim 0.1 \mu\text{M}$ .

5

### 6 **3.2. Biomarker lipid profiles**

7 The SPM for biomarker analysis were collected on  $0.7 \mu\text{m}$  pore size GF filters. Limitations  
8 related to the use of  $0.7 \mu\text{m}$  filters to collect archaeal living cells have been reported (Ingalls  
9 et al., 2012; Schouten et al., 2012) as the typical size of thaumarchaeotal cells is  $<0.6 \mu\text{m}$   
10 (Könneke et al., 2005) and they are suggested to occur predominantly free-living during their  
11 lifetime (Ingalls et al., 2012). Although the pore size tends to diminish as the particulate  
12 material accumulates, the employment of  $0.7 \mu\text{m}$  filters likely causes an underestimation of  
13 the archaeal population, and thus archaeal IPL abundance (Schouten et al., 2012). However,  
14 Pitcher et al. (2011b) showed that depth profiles of HPH-crenarchaeol, analyzed on SPM  
15 collected using  $0.7 \mu\text{m}$  GFF filters, in the Arabian Sea ODZ were similar to that of  
16 thaumarchaeotal genes, analyzed on SPM collected using  $0.2 \mu\text{m}$  filters (Pitcher et al.,  
17 2011b). Therefore, our results are likely still suitable to probe the depth habitat of  
18 Thaumarchaeota.

19 Figures 3e and 3m show HPH-crenarchaeol vertical profile for the coastal and the  
20 open ocean sites, respectively. In the coastal setting, HPH-crenarchaeol has a maximum in  
21 abundance at the interface between the oxycline and the upper ODZ, where  $\sigma_\theta$  is  $\sim 25$  (i.e.  $\sim 30$   
22 m depth) (Fig. 3e). In the open ocean setting, HPH-crenarchaeol starts to increase at declining  
23 oxygen concentrations and peaks at the base of the oxycline (i.e. at  $\sigma_\theta$  25.8 and 100 m depth).  
24 Deeper in the water column (i.e. at  $\sigma_\theta$  27.4 corresponding to  $\sim 1000$  m water depth) a  
25 secondary minor maximum in HPH-crenarchaeol was detected (Fig. 3m).

1 In the coastal waters, the ladderane PC-monoether concentration stays low, except for  
2 one data point (i.e. ~17 pg L<sup>-1</sup> at 55 m depth), until the upper ODZ where it starts to increase  
3 to its maximum (i.e. ~251 pg L<sup>-1</sup>) at  $\sigma_{\theta}$  26.4 in the core ODZ (i.e. 150 m depth) (Fig. 3f). In  
4 the open ocean, the PC-monoether maximum in concentration (i.e. ~122 pg L<sup>-1</sup>) is located at  
5 the oxycline (i.e. at  $\sigma_{\theta}$  25.9 and 105 m depth).

## 6 7 **4. Discussion**

### 8 **4.1. Depth distributions and abundance of AOA and anammox bacteria in** 9 **the ETNP**

10 In this study we have been able to investigate concurrently for the first time, the vertical  
11 distribution of AOA and anammox bacteria in both coastal and open waters of the ETNP  
12 ODZ. The IPL-biomarker profiles show that AOA and anammox bacteria are present in the  
13 region and partially co-exist along the water column (Figs. 3e and 3f; 3m and 3n). Such a  
14 distribution was already observed in other dysoxic or anoxic marine systems worldwide such  
15 as the Black Sea, and the ODZs of the ESTP and the Benguela upwelling system (Lam et al.,  
16 2007, 2009; Woebken et al., 2007), whereas in the southern part of the ETNP ODZ (Podlaska  
17 et al., 2012) and in the Arabian Sea (Pitcher et al., 2011b) the two microbial groups are  
18 reported to thrive at different water depths. In the northern ETNP our IPLs depth profiles  
19 highlight some substantial differences in the distribution and abundance of the two groups  
20 between the different settings.

21 In the coastal setting, the two microbial groups show clear niche segregation in the  
22 upper part of the water column. Here, AOA thrive at the bottom of the oxycline, at a  $\sigma_{\theta}$  of  
23 ~25, whereas anammox bacteria are just starting to increase in abundance at that point and  
24 exhibit a clear maximum only in the core ODZ (Fig. 3e and 3f), where  $\sigma_{\theta}$  has shifted to ~26.  
25 The trend of our coastal HPH-crenarchaeol depth profile agrees with previously reported data

1 for thaumarchaeotal 16SrRNA, archaeal *amoA* gene concentration and rate measurements  
2 from the same area (station 3 in Beman et al., 2012), which also revealed an AOA maximum  
3 at the base of the oxycline. Consequently, Beman et al. (2012) suggested a prominent role of  
4 AOA in performing nitrification in shallow O<sub>2</sub>-depleted waters. The observed maximum  
5 abundance of anammox bacteria in the core ODZ as based on the ladderane lipid profile is in  
6 agreement with previous investigations in the ETSP ODZ, where it has been proposed as a  
7 preferential niche for anammox activity (De Brabandere et al., 2014; Hamersley et al., 2007;  
8 Ward et al., 2009). Moreover, similarly to De Brabandere et al. (2014), who also reported low  
9 anammox rates at the oxycline in one of their sampling stations in the ETSP, we also observed  
10 low ladderane concentrations in the ETNP coastal setting.

11 At the open ocean site, we also find the maximum abundance of anammox bacteria  
12 between the base of the oxycline and the upper part of the ODZ. However, here the anammox  
13 bacterial abundance displays a concurrent maximum with that of AOA (Figs. 3m and 3n). The  
14 segregation of AOA and anammox bacteria niches in the coastal waters of the ETNP ODZ  
15 and their contrasting co-occurrence in the open waters clearly indicates a different behavior of  
16 the two microbial species at different locations of the ETNP. To the best of our knowledge  
17 this is the first study that highlights such different vertical distribution of the two groups on a  
18 local scale. We also note that both IPL-biomarkers exhibit higher concentration maxima in  
19 coastal waters (Figs. 3e and 3f) than in the open ocean (Figs. 3m and 3n), i.e. the  
20 concentration of HPH-crenarchaeol is five times higher and that of the PC-monoether  
21 ladderane is more than twice that found in the open ocean. This suggests that both AOA and  
22 anammox bacteria are more abundant in the coastal waters of the ETNP. The reasons for such  
23 divergence may be various. For instance, the complex and so far not fully resolved upper  
24 water circulation in this region may play a role (Fiedler and Talley, 2006; Kessler, 2006). The  
25 proximity of the American continent is likely to have a greater influence on the hydrography

1 of the coastal site (i.e. in a straight line the closest point on the Mexican coastline to our  
2 coastal settings is roughly 40 km away), than on the open ocean site. At these latitudes the  
3 continental wind forcing is a dominant factor and together with the variations in the coastline  
4 influences the local upper circulation (Fiedler and Talley, 2006; Kessler, 2006) and might  
5 have an effect on the different vertical distribution and abundance observed in the two  
6 microbial species as well. In the same way the nitrogen species profiles are likely to be  
7 influenced by variable hydrographical features (Fig. 3).

8

#### 9 **4.2. Influence of nitrogen species on the abundance and the distribution of** 10 **Thaumarchaeota and anammox bacteria in the ETNP**

11 Ammonium and nitrite concentrations have been proposed as critical factors in determining  
12 the vertical distribution and the abundance of Thaumarchaeota and anammox bacteria  
13 (Hamersley et al., 2007; Jaeschke et al., 2007; Jensen et al., 2009; Kuypers et al., 2005;  
14 Martens-Habbena et al., 2009; Stahl and de la Torre, 2012; Thamdrup et al., 2006; Ward et  
15 al., 2009).

16  $\text{NH}_4^+$  serves as a substrate for both Thaumarchaeota and anammox bacteria and has  
17 been observed to not accumulate in ODZs as a results of efficient turnover between sources  
18 and sinks (Kalvelage et al., 2013). In the ETNP ODZ, we found both Thaumarchaeota and  
19 anammox bacteria in sub-oxic and anoxic waters. Ammonium concentrations are low and  
20 mostly under the detection limit, likely due to the consumption of the nutrient by both (or  
21 other) microorganisms (Figs. 3c and 3i). Even at concentrations  $<1 \mu\text{M}$ ,  $\text{NH}_4^+$  may support  
22 anammox reaction, which is considered the main sink for this nitrogen species in the core  
23 ODZs (Bianchi et al., 2014). Nitrite is the electron acceptor in the anammox process and it has  
24 been already described as a limiting factor to anammox bacteria activity in the southern ETNP  
25 ODZ (Rush et al., 2012).



1           In the coastal waters, thaumarchaeotal nitrification is probably taking place at the  
2 bottom of the oxycline, as indicated by the HPH-crenarchaeol maximum (Fig. 3e) and the  
3 concurrent ammonium concentration peak (Fig. 3c), most likely resulting from the  
4 mineralization of organic matter. Moreover, thaumarchaeotal nitrification, which converts  
5 ammonium into nitrite (Arrigo, 2005), may cause the observed minor primary peak in the  
6 nitrite concentration profile (the so-called primary nitrite maximum or PNM) occurring at the  
7 bottom of the oxycline in these waters (Fig. 3b), which coincides with the maximum of AOA  
8 abundance (Figs. 3b and 3e). In the core ODZ a clear secondary nitrite maximum (SNM) co-  
9 occurs with the maximum in anammox bacteria concentration (Figs. 3b and 3f). Although  
10 heterotrophic denitrification represents the obvious candidate as main provider of nitrite to  
11 anammox bacteria in oceanic settings (Ward et al., 2009), the two processes are usually not  
12 found coupled together (Dalsgaard et al., 2012). Alternatively, a combination of several  
13 pathways including dissimilatory  $\text{NO}_3^-$  reduction to  $\text{NO}_2^-$  (DNRN) or  $\text{NH}_4^+$  (DNRA) plays a  
14 role, as has been observed in other ODZs (Canfield et al., 2010; Kartal et al., 2007; Lam et al.,  
15 2009, 2011; Lipschultz et al., 1990; Ward et al., 2009). The extent of the contribution of these  
16 processes as nutrients providers to anammox bacteria is still unclear (Lam and Kuypers,  
17 2011). Finally, a recent study has also brought into attention zooplankton migrators as  
18 alternative source of substrates to the anammox metabolism, previously overlooked in ODZs  
19 (Bianchi et al., 2014). In total, these things suggest that those mechanisms are all feasible to  
20 feed the anammox process in the coastal waters of the core ETNP ODZ with the nutrients  
21 required.

22           In the open ocean ETNP, AOA and anammox bacteria maxima may be coupled in the  
23 upper ODZ because of the (partial) overlap of the ecological niches of the two groups in this  
24 setting (Figs. 3m and 3n). In this case, AOA and anammox bacteria could either compete or  
25 co-operate for the substrates they require, as already proved in laboratory-scale models (Yan

1 et al., 2010, 2012). The nutrient profiles are also consistent with the co-occurrence of the two  
2 metabolic processes (Figs. 3h and 3i): the secondary nitrite maximum is concurrent with the  
3 two biomarker maxima (Figs. 3h; 3m and 3n), whereas  $\text{NH}_4^+$  is either possibly consumed by  
4 anammox as the water column turns sub-oxic or by nitrification by AOA successfully adapted  
5 to low nutrients and oxygen conditions (Coolen et al., 2007; Francis et al., 2005; Lam et al.,  
6 2007, 2009; Martens-Habbena et al., 2009; Park et al., 2010; Pitcher et al., 2011b; Schouten et  
7 al., 2004; Sinninghe Damsté et al., 2002a; Stahl and de la Torre, 2012; Stolper et al., 2010;  
8 Tiano et al., 2014; Woebken et al., 2007) and potentially to a broad variety of different  
9 environmental conditions (Qin et al., 2014).

10 The role of  $\text{NO}_2^-$  and  $\text{NH}_4^+$  in differentiating the distribution of Thaumarchaeota and  
11 anammox bacteria observed at the coastal and at the open water sites of the ETNP ODZ is not  
12 clear; when the  $\text{NO}_2^-$  and  $\text{NH}_4^+$  concentrations were compared with those of the specific  
13 biomarkers studied at both sites, no evident relationship was apparent (data not shown).

14 In conclusion, further investigation is required to establish the contribution of the  
15 single processes to the N cycle occurring in the settings investigated in this study and to  
16 explain the divergence between the two. Other studies have called attention to the relevance  
17 of organic matter fluxes as a control over these metabolic pathways and ultimately over the  
18 balance between the two mainly responsible for the  $\text{N}_2$  removal from the oceans, i.e.  
19 anammox and denitrification (Babbin et al., 2014; Chang et al., 2014; Kalvelage et al., 2013;  
20 Koeve and Kähler, 2010; Ward, 2013; Ward et al., 2008, 2009). Specifically, variations in the  
21 C/N ratio content of the particulate organic matter (POM) entering the ODZ may play a  
22 prominent role in determining anammox and heterotrophic denitrification rates, with  
23 anammox being favorite by nitrogen-rich OM (Babbin et al., 2014).

24

#### 25 **4.3. The role of the oxygen**

1 As the features of ODZs suggest, oxygen might play a pivotal role in controlling the  
2 abundance and the special distribution of Thaumarchaeota and anammox bacteria in those  
3 areas. Previous studies have already pointed to this in other ODZs (Jaeschke et al., 2007;  
4 Kuypers et al., 2005; Stahl and de la Torre, 2012; Thamdrup et al., 2006) and in the southern  
5 ETNP ODZ itself (Rush et al., 2012). To investigate if oxygen concentration is influencing  
6 the abundance of Thaumarchaeota and anammox bacteria in our study sites in the ETNP  
7 ODZ, we compared our biomarker concentrations with oxygen concentrations in both coastal  
8 and open ocean site (Figs. 4a and 4b). Figure 4a and 4b shows how PC-monoether ladderane  
9 and HPH-crenarchaeol are distributed according to O<sub>2</sub> concentration at the two sites. The two  
10 distributions appear rather similar, with both biomarkers being more abundant at an oxygen  
11 concentration below the detection limit, i.e. ca. 1 μM, which is even overestimated by the  
12 CTD sensor employed for the measurements, as suggested by O<sub>2</sub> measurements taken with  
13 the STOX microsensor during the same cruise (Tiano et al., 2014). The only evident  
14 difference between the two plots is found in one HPH-crenarchaeol data point from the  
15 coastal site, corresponding to an oxygen concentration of ~26 μM, which reflects the much  
16 broader range of tolerance of AOA to O<sub>2</sub> compared to the strictly anaerobic anammox  
17 bacteria (Tiano et al., 2014). However, the relation revealed by our plots suggests that both  
18 microbial species are potentially able to cope with low oxygen concentrations and O<sub>2</sub> plays a  
19 primary role in controlling the distribution of the two microbial species, as shown previously  
20 (Martens-Habbena et al., 2009; Park et al., 2010; Pitcher et al., 2011b; Rush et al., 2012; Stahl  
21 and de la Torre, 2012; Tiano et al., 2014). The high relative abundance of HPH-crenarchaeol  
22 in the poorly oxygenated waters of the ETNP is consistent with the ability of AOA to thrive  
23 and perform nitrification under low oxygen conditions (Coolen et al., 2007; Francis et al.,  
24 2005; Lam et al., 2007, 2009; Park et al., 2010; Pitcher et al., 2011b; Schouten et al., 2004;  
25 Sinninghe Damsté et al., 2002a; Stolper et al., 2010; Woebken et al., 2007). In the open ocean

1 site a secondary minor peak of HPH-crenarchaeol at the lower oxycline, i.e. 1100 m depth  
2 (Fig. 3m), supports the hypothesis. Pitcher et al. (2011b) also observed a secondary maximum  
3 of AOA at the bottom of the Arabian Sea ODZ. Our findings thus confirm oxygen  
4 concentration as an important environmental control in determining the distribution of  
5 Thaumarchaeota and anammox bacteria in the water column of the ETNP ODZ.

6

#### 7 **4.4. Conclusions**

8 In this study high resolution profiles of the two specific IPL-biomarkers of AOA and  
9 anammox bacteria, i.e. HPH-crenarchaeol and PC-monoether ladderane, allowed us to have a  
10 detailed insight on the vertical distribution of these microbial groups in the ETNP ODZ. It  
11 shows that AOA and anammox bacteria are abundant at both shallow coastal and open ocean  
12 waters of the ETNP oxygen deficient zone. Our findings also indicate that the ecological  
13 niches of the two species diverge on a local scale in the ETNP. Different O<sub>2</sub> concentration and  
14 water stratification features between the two study sites play an important role in determining  
15 such differences; whereas the role of NO<sub>2</sub><sup>-</sup> and NH<sub>4</sub><sup>+</sup> is not clear. Further studies are needed to  
16 elucidate potential interactions between AOA and anammox in this ODZ. However future  
17 investigations on the N-cycle in the ETNP and other ODZs might take into a greater account  
18 the importance of regional differences in the ecological niches of these microbial species.

19

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39 Table 1. Sampling locations for SPM during the Eastern Tropical North Pacific cruise aboard the R/V Thomas G. Thompson (March-April  
 40 2012). For each sampling location the table reports the depth of SPM sampling, the volume of water filtered by each pump deployed as well as  
 41 physical parameters of the water at these depths, i.e. temperature and oxygen concentration (O<sub>2</sub>). Nutrients concentrations, i.e. nitrate (NO<sub>3</sub><sup>-</sup>),  
 42 nitrite (NO<sub>2</sub><sup>-</sup>) and ammonium (NH<sub>4</sub><sup>+</sup>) and corresponding station and depth of sampling, when available, are also reported.

| Station    | Location      | Temperature<br>CTD | Depth of<br>SPM<br>sampling | Water<br>filtered | O <sub>2</sub> | Depth of<br>nutrients<br>sampling | NO <sub>3</sub> <sup>-</sup> | NO <sub>2</sub> <sup>-</sup> | NH <sub>4</sub> <sup>+</sup> |
|------------|---------------|--------------------|-----------------------------|-------------------|----------------|-----------------------------------|------------------------------|------------------------------|------------------------------|
|            |               | (°C)               | (m)                         | (L)               | (µM)           | (m)                               | (µM)                         | (µM)                         | (µM)                         |
| <b>106</b> | 20°14.49' N   | 15                 | 70                          | 1627.72           | 0.75           | n.d.                              | n.d.                         | n.d.                         | n.d.                         |
|            | 106°10.00' W  | 13                 | 105                         | 200.00            | 0.81           | n.d.                              | n.d.                         | n.d.                         | n.d.                         |
|            |               | 10                 | 365                         | 699.50            | 0.97           | n.d.                              | n.d.                         | n.d.                         | n.d.                         |
| <b>110</b> | 20° 08.48' N  | 14                 | 70                          | 128.50            | 0.67           | 50                                | 20.14                        | 2.32                         | 0.02                         |
|            | 105° 59.18' W | 13                 | 125                         | 336.14            | 0.77           | 100                               | 15.27                        | 7.06                         | 0.01                         |
|            |               | 12                 | 150                         | 511.79            | 0.86           | 150                               | 20.13                        | 5.12                         | 0.00                         |
|            |               | 6                  | 710                         | 999.50            | 1.05           | n.d.                              | n.d.                         | n.d.                         | n.d.                         |
| <b>114</b> | 20° 07.54' N  | 18                 | 25                          | 104.10            | 26.20          | n.d.                              | n.d.                         | n.d.                         | n.d.                         |
|            | 106° 00.84' W | 17                 | 35                          | 97.40             | 0.78           | n.d.                              | n.d.                         | n.d.                         | n.d.                         |
|            |               | 15                 | 45                          | 255.51            | 0.71           | n.d.                              | n.d.                         | n.d.                         | n.d.                         |
|            |               | 15                 | 55                          | 230.50            | 0.71           | n.d.                              | n.d.                         | n.d.                         | n.d.                         |
| <b>119</b> | 20° 08.72' N  | 14                 | 80                          | 211.60            | 0.72           | 100                               | 16.55                        | 5.74                         | 0.00                         |
|            | 105° 59.77' W | 8                  | 500                         | 932.50            | 0.95           | 501                               | 31.27                        | 1.50                         | 0.00                         |
|            |               | 5                  | 800                         | 595.50            | 1.61           | 800                               | 43.66                        | 0.00                         | 0.00                         |
|            |               | 5                  | 1000                        | 1069.38           | 8.78           | 1001                              | 45.42                        | 0.00                         | 0.00                         |

|                |               |      |                   |         |        |      |       |      |      |
|----------------|---------------|------|-------------------|---------|--------|------|-------|------|------|
| <b>123</b>     | 20° 03.50' N  | 19   | 20                | 41.50   | 51.16  | n.d. | n.d.  | n.d. | n.d. |
|                | 106° 00.59' W | 14   | 60                | 181.70  | 0.71   | n.d. | n.d.  | n.d. | n.d. |
|                |               | 13   | 90                | 179.81  | 0.76   | n.d. | n.d.  | n.d. | n.d. |
|                |               | 12   | 200               | 210.50  | 0.87   | n.d. | n.d.  | n.d. | n.d. |
| <b>125-126</b> | 20° 04.26' N  | 20   | 15 <sup>a</sup>   |         | 72.05  | 15   | 12.55 | 0.96 | 0.36 |
|                | 106° 00.81' W | n.d. | 1300              | 1003.00 | n.d.   | n.d. | n.d.  | n.d. | n.d. |
|                |               | n.d. | 1600 <sup>a</sup> |         | n.d.   | n.d. | n.d.  | n.d. | n.d. |
|                |               | n.d. | 2000              | 859.66  | n.d.   | n.d. | n.d.  | n.d. | n.d. |
| <b>136</b>     | 17° 01.95' N  | 14   | 110               | 755.19  | 0.74   | 110  | 21.54 | 1.07 | 0.00 |
|                | 106° 31.96' W | 13   | 150               | 747.50  | 0.74   | 160  | 19.24 | 4.74 | 0.00 |
|                |               | 11   | 250               | 987.99  | 0.87   | 200  | 21.30 | 3.37 | 0.00 |
|                |               | 10   | 350               | 714.90  | 0.88   | 300  | 23.39 | 1.97 | 0.00 |
| <b>141</b>     | 16° 30.98' N  | 23   | 60                | 482.64  | 108.13 | 60   | 7.00  | 1.14 | 0.16 |
|                | 107° 08.52' W | 16   | 90                | 997.00  | 0.88   | 80   | 22.91 | 0.45 | 0.00 |
|                |               | 15   | 105               | 496.00  | 0.77   | 100  | 24.20 | 0.09 | 0.00 |
|                |               | 12   | 200               | 864.96  | 0.89   | 181  | 23.62 | 5.21 | 0.00 |
| <b>145</b>     | 16° 31.78' N  | 26   | 50                | 368.00  | 189.01 | n.d. | n.d.  | n.d. | n.d. |
|                | 107° 08.45' W | 21   | 65 <sup>a</sup>   |         | 69.83  | n.d. | n.d.  | n.d. | n.d. |
|                |               | 13   | 155               | 768.44  | 0.99   | n.d. | n.d.  | n.d. | n.d. |
|                |               | n.d. | 710               | 1417.63 | n.d.   | n.d. | n.d.  | n.d. | n.d. |
| <b>147-149</b> | 16° 31.60' N  | 13   | 170               | 668.00  | 0.84   | 175  | 27.54 | 4.04 | 0.00 |
|                | 107° 06.80' W | 5    | 990               | 1249.18 | 7.30   | n.d. | n.d.  | n.d. | n.d. |
|                |               | 4    | 1100              | 689.50  | 13.11  | 1100 | 46.27 | 0.01 | 0.00 |
|                |               | n.d. | 2500              | 567.81  | n.d.   | n.d. | n.d.  | n.d. | n.d. |
| <b>154-155</b> | 16° 35.34' N  | 25   | 55                | 382.80  | 176.12 | n.d. | n.d.  | n.d. | n.d. |

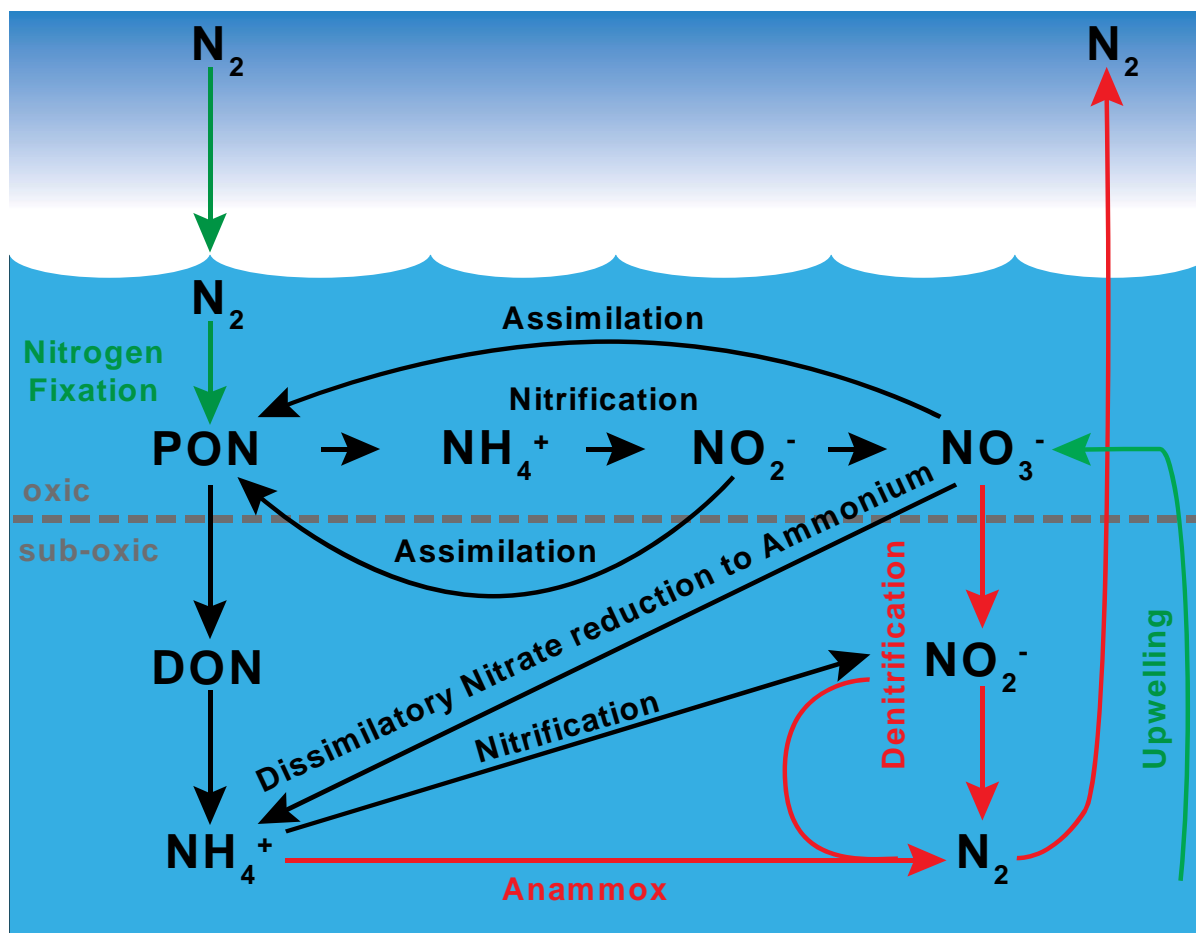
|                |               |      |                  |         |      |      |       |      |      |
|----------------|---------------|------|------------------|---------|------|------|-------|------|------|
|                | 107° 08.98' W | 15   | 100              | 686.29  | 1.09 | 104  | 23.40 | 0.65 | 0.00 |
|                |               | 13   | 145              | 334.50  | 1.06 | n.d. | n.d.  | n.d. | n.d. |
|                |               | n.d. | 160              | 612.86  | n.d. | n.d. | n.d.  | n.d. | n.d. |
| <b>157-158</b> | 16° 34.66' N  | 19   | 80               | 3287.62 | 3.49 | n.d. | n.d.  | n.d. | n.d. |
|                | 107° 04.61' W | 13   | 140              | 1178.02 | 0.75 | 141  | 25.70 | 3.89 | 0.02 |
|                |               | 9    | 450              | 3484.84 | 0.96 | n.d. | n.d.  | n.d. | n.d. |
|                |               | 8    | 550 <sup>a</sup> |         | 1.02 | n.d. | n.d.  | n.d. | n.d. |

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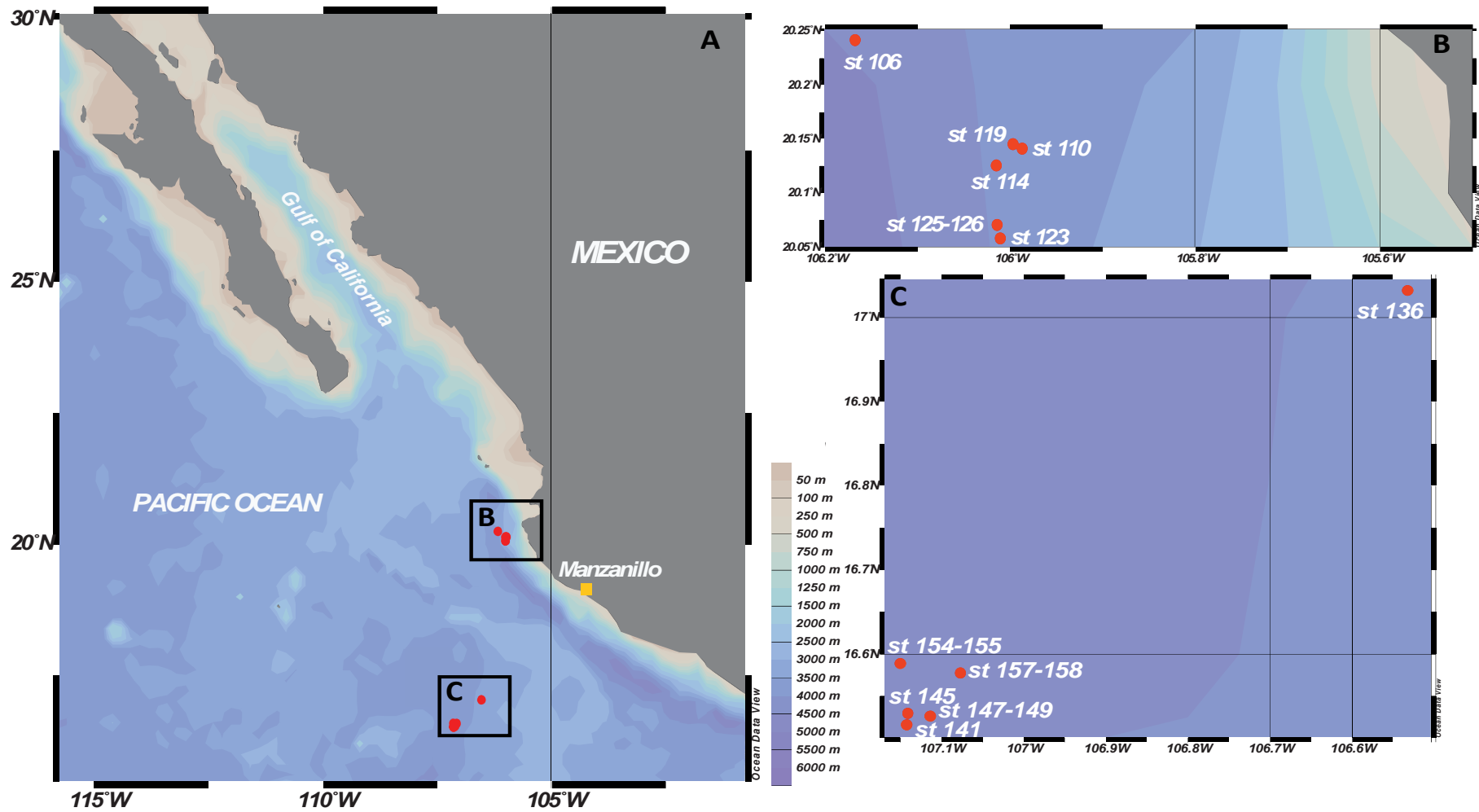
44 <sup>a</sup> Sample not analyzed for lipids.





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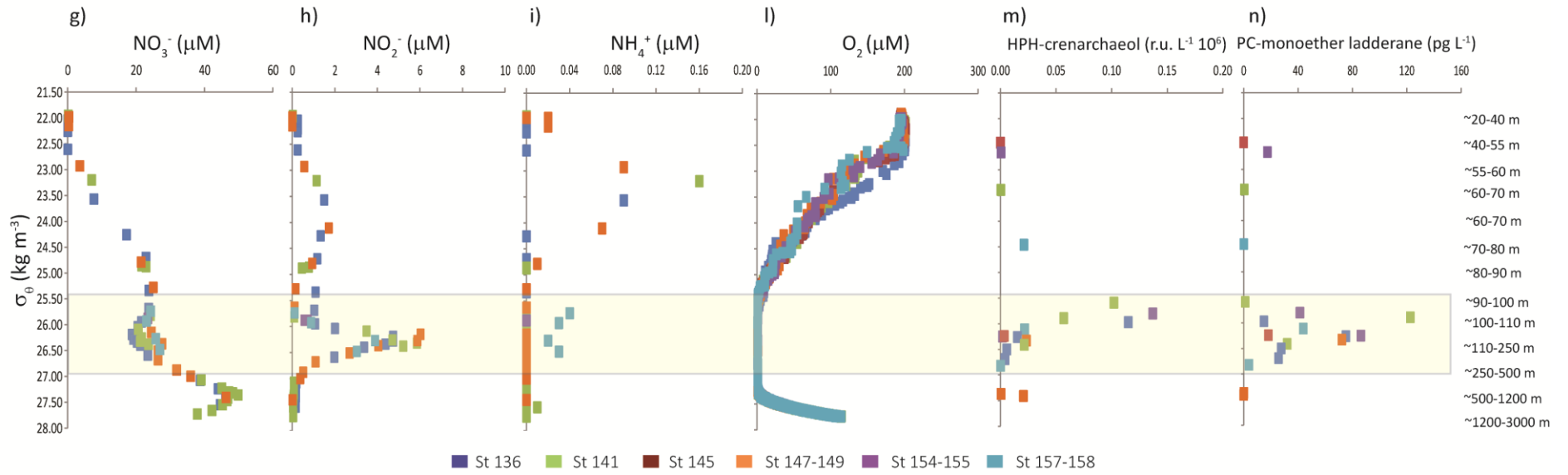
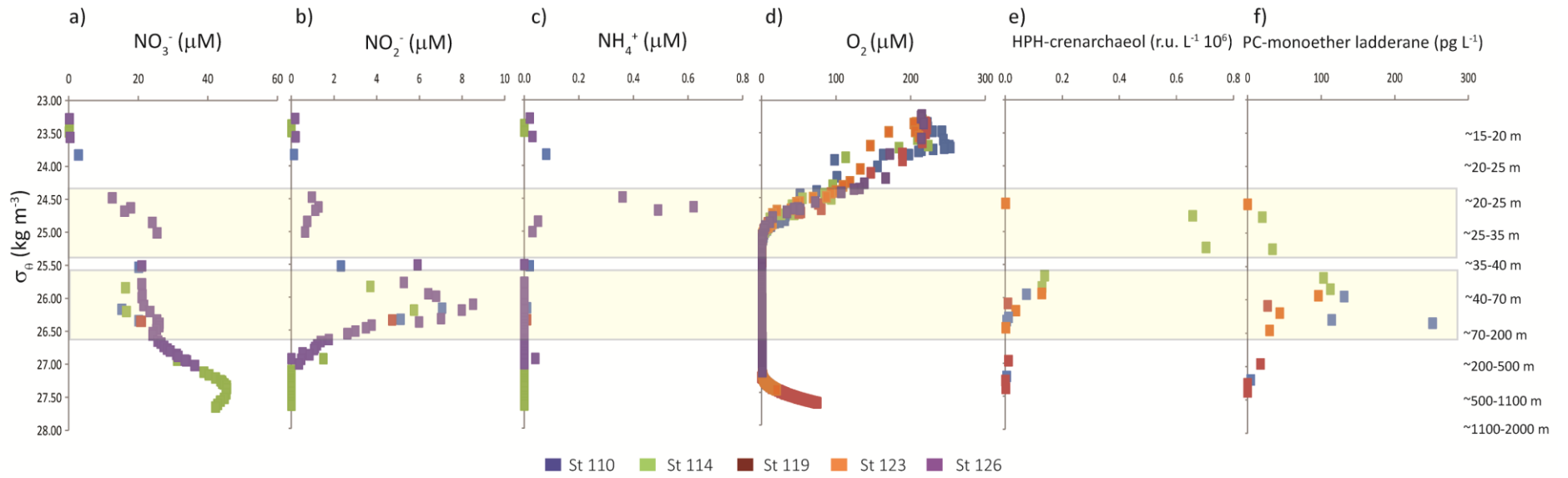
46 Figure 1. The marine biogeochemical nitrogen cycle. The main redox reactions involved are included: in green are the processes responsible for  
 47 gains of nitrogen to the ocean, in red are the losses. Modified from Arrigo (2005).



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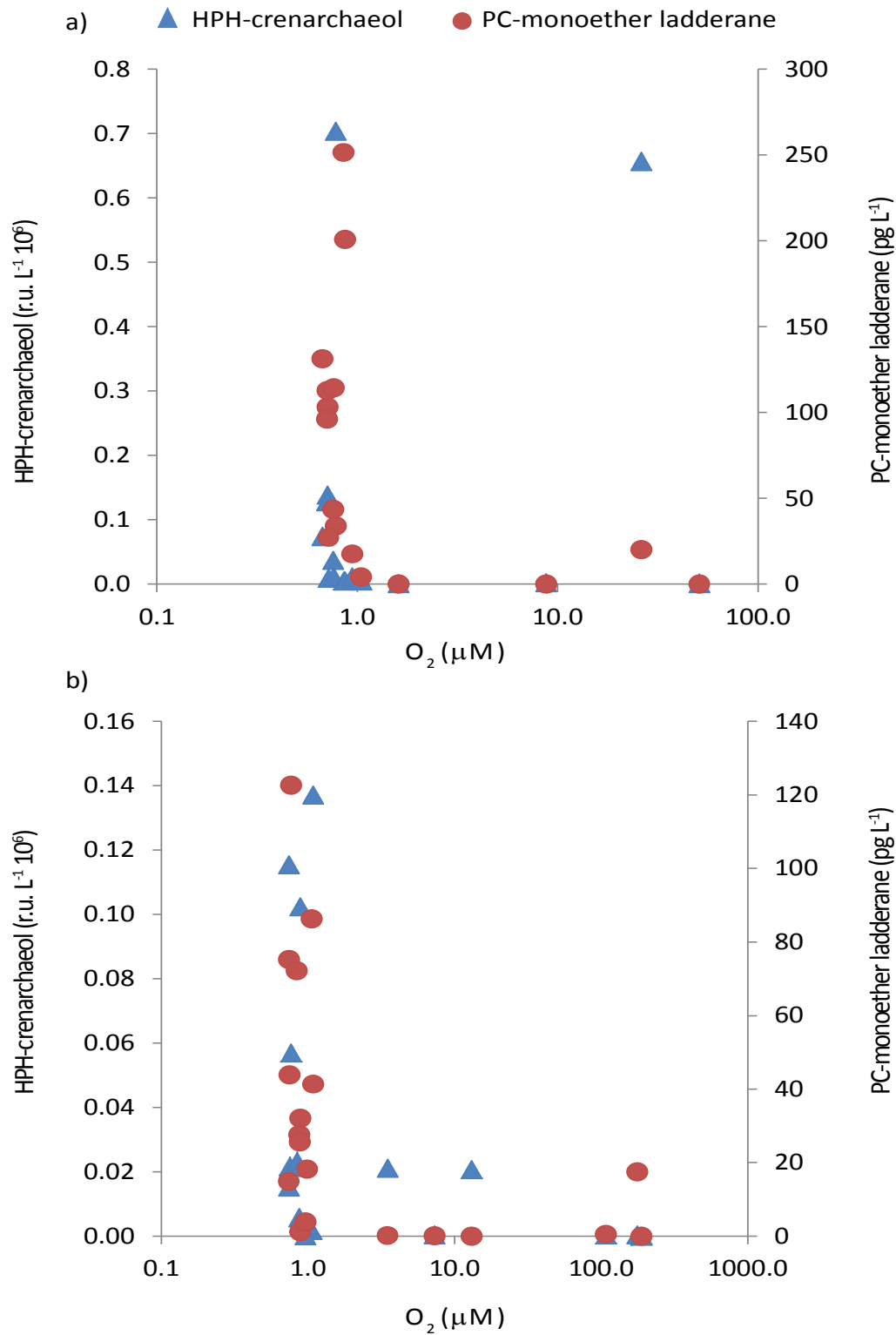
49 Figure 2. (A) Map of the sampling area of the Eastern Tropical North Pacific cruise (March-April 2012) and sampling stations (red dots), in the  
 50 coastal (B) and in the open ocean site (C). The coastal sampling site (i.e. st 106, st 110, st 114, st 119, st 123, and st 125-126) is placed in the area

- 51 between 20 25.00N and 105 60.00W; the open ocean sampling site (i.e. st 136, st 141, st 145, st 147-149, st 154-155, st 157-158) is placed in the
- 52 area between 17 00.00N and 106 60.00W.



54 Figure 3. Concentration profiles of **(a, g)** nitrate ( $\text{NO}_3^-$ ); **(b, h)** nitrite ( $\text{NO}_2^-$ ); **(c, i)** ammonium ( $\text{NH}_4^+$ ); **(d, l)** oxygen; **(e, m)** hexose-  
55 phosphohexose (HPH)-crenarchaeol; and **(f, n)** phosphocholine (PC)-monoether ladderane according to the potential density anomalies ( $\sigma_\theta$ ) of  
56 the water column of the ETNP. The corresponding depths intervals are reported on the right side of the figure as a reference. Upper four panels  
57 **(a-f)** provide an overview of the complete water column (2000 m) of the coastal sampling site; lower four panels **(g-n)** show the complete water  
58 column (3000 m) relative to the open ocean sampling site. All profiles are obtained by combining respectively the coastal and the open ocean  
59 stations sampled. The yellow bars highlight the main differences observed between the two sampling sites with respect to the distribution of the  
60 two microbial species: i.e. on the one hand a clear niche segregation of AOA and anammox bacteria in the coastal waters of the ETNP and on the  
61 other hand a partial overlap of the two niches of these microbial species in the open water setting.

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63

64 Figure 4. Abundance of HPH-crenarchaeol ( $\text{r.u. L}^{-1}$ , i.e. response units  $\text{L}^{-1}$ ) and PC-monoether  
 65 ladderane lipids ( $\text{pg L}^{-1}$ ) according to the concentration of oxygen ( $\mu\text{M}$ ). In the upper panel  
 66 (a) the response of the two biomarkers at the coastal site; in the lower panel (b) the response  
 67 at the open ocean site.