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- The successional formation and release of domoic acid in a *Pseudo-nitzschia* bloom in
 the Juan de Fuca Eddy: a drifter study.
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- 20 Abstract
- 21

22 Blooms of Pseudo-nitzschia species are frequent, but presently unpredictable, in the 23 Juan de Fuca Eddy region off the coasts of Washington (US) and British Columbia 24 (Canada). This upwelling eddy region is proposed to be the bloom initiation site, from 25 nutrients upwelled to the surface, before cells are entrained into the coastal surface 26 currents. During a shipboard study, we characterized the different stages of the *Pseudo*-27 nitzschia bloom development from its initiation and intensification, to its eventual 28 sinking and dissipation. Specifically, we followed a water mass using lagrangian 29 ARGOS-tracked drifters released at the eddy water mass and quantified production of 30 dissolved and particulate domoic acid, and the physiological status of the Pseudo-31 *nitzschia* cells with regards to photosynthesis, nutrient needs and sinking rates, along 32 with its relationship with competing species – in this case, the marine euglenoid, 33 *Eutreptiella* spp. The drifter study allows for an interpretation of the presence or 34 absence of Pseudo-nitzschia and domoic acid against active environmental factors -35 particularly copper and iron. 36

Key Words: algal biotoxins, amnesic shellfish poisoning, copper, DFB, domoic acid,
Harmful Algal Blooms, iron, Juan de Fuca Eddy, *Pseudo-nitzschia*, Washington State

1. Introduction

43 The Juan de Fuca eddy is a nutrient-rich, physically retentive region off the 44 northwestern U.S. coast that serves as an initiation site for toxigenic diatoms of the 45 Pseudo-nitzschia genus Pergallo [Heterokonta, Bacillariophyceae] (Trainer et al., 2002; 46 Trainer and Hickey, 2009; Hickey et al., 2013). This region serves as a natural 47 laboratory for studying the *Pseudo-nitzschia* cells, and the oceanographic progression of 48 the blooms at the eddy core, where nutrients are introduced, to the possible intersection 49 of the bloom to the coastline (Adams et al., 2000, Trainer et al., 2002), where domoic 50 acid is introduced to economically, socially, and culturally significant shellfish 51 populations (Chadsey et al., 2011; Dyson and Huppert, 2010). Less studied are the 52 physiological changes associated with the progression of *Pseudo-nitzschia* from the 53 bloom initiation site to the presumably more mature bloom some distance and time 54 away from the initiation site.

55 Historically, researchers have used a spatial census method to systematically acquire 56 and record data about the members of a given bloom-forming population. This approach 57 is valuable, links well with standardized surveys for physical and chemical profiles and 58 provides a spatial template from which the trajectory and path of the bloom cells can be 59 transcribed. The alternative is to plan the research and cruise track "to follow the 60 bloom", or rather, to follow a drifter over time that is designed to track water at a 61 specific depth – water that contains the planktonic bloom cells under study. This 62 Lagrangian approach is comparable to a population cohort survey, where the same 63 population is followed over time, and key population factors such as growth, loss, photosynthetic efficiency and physiological health can be assessed within the bloom 64

65 (Lewitus et al., 2012; Trainer et al., 2012a). This latter approach enables measurements 66 at one point in time to be confidently connected with previous and future measurements 67 The Ecology and Oceanography of Harmful Algal Blooms in the Pacific Northwest 68 (ECOHAB-PNW) was a 5-year, 6-cruise multi-disciplinary project that measured 69 physical, chemical, and biological parameters associated with the periodic blooms of 70 Pseudo-nitzschia species off the coasts of Washington State and Vancouver Island, 71 British Columbia including the entrance waters of the Strait of Juan de Fuca. In 72 September 2004, the cruise strategy included the release of Lagrangian ARGOS-tracked 73 drifters that when placed at the sources of nutrients (eddy core; Hickey et al., 2006; 74 MacPhadyen et al., 2008) enabled the longitudinal study of the phytoplankton 75 community. The multi-week cruise occurred during a large, nearly monospecific diatom 76 bloom of *P. cuspidata* that co-dominated the phytoplankton assemblage with the 77 euglenoid, Eutreptiella spp. (Trainer et al., 2009b). There were no significant 78 correlations between the observed domoic acid (DA) concentrations [particulate DA 79 (pDA) and cellular DA] and the ambient concentrations of macronutrients (nitrate, 80 orthophosphate and silicate) (Trainer et al. 2009b). As with the majority of the spatial 81 census cruises, neither were correlations detected between pDA or *Pseudo-nitzschia* 82 concentrations and total bacteria or cyanobacteria abundances. Combined with the 83 correlational assessment of environmental, ecological and toxicological data, we used a 84 series of grow-out experiments to consider if the postulate that domoic acid acted as a 85 ligand to bind either copper or iron (Rue and Bruland, 2001), improving the 86 physiological health of the cells as these putatively limiting trace metals were 87 scavenged from the environment (Maldonado et al., 2002; Wells et al., 2005), enabling 88 Pseudo-nitzschia to achieve less restricted physiological health Here we demonstrate

how the independent nature of *Pseudo-nitzschia* is best revealed by the variation in two
trace metals – copper and iron – working in conjunction with particulate and dissolved
DA levels.

92

93 **2. Materials and methods**

94 2.1. Cruise

The ECOHAB-PNW-III cruise was carried out aboard the R/V *Atlantis* (AT1117) during 8-28 September 2004. The initial survey region for a large-scale synoptic
assessment covered *ca*. 12,000 km⁻², spanning a N-S latitude line from 48.7°N to 47.0
°N, along a longitude of 125.0°W. The synoptic scale survey grid was designed to
include areas influenced by the Strait of Juan de Fuca, the Juan de Fuca Eddy region
and the coastal upwelling region off the Washington coast.

102 2.2. Drifters

Lagrangian ARGOS-tracked drifters (Brightwater Instrument Co. models 104a and 115) were deployed to delineate patterns and speeds of surface flows in the eddy area, as well as to determine the ultimate fate of eddy water. These drifter models were designed according to the Davis/CODE configuration to accurately track the upper 1 m of the water column (Davis, 1985), and transmitted ½ hourly GPS position to the ARGOS satellites. Drifter A was deployed in the outer eddy at 21:00 h on September 16 and was followed for 10 days.

110

111 2.3. Satellite imagery

112	Sea-viewing wide field-of-view sensor (SeaWiFS) imagery was acquired from
113	the National Oceanic and Atmospheric Administration's (NOAA) Coastwatch Program.
114	The images were processed with the latest version of SeaWiFS data analysis system
115	(SeaDAS 4.0), which uses an atmospheric correction that compensates for near-infrared
116	water leaving radiance and absorbing aerosols (Gordon and Wang, 1994; Stumpf et al.,
117	2003). The resulting chlorophyll imagery was developed using the global OC4
118	algorithm, with 1 km resolution (O'Reilly et al., 2000).
119	
120	2.4. Chlorophyll a
121	Surface samples were analyzed for phytoplankton biomass as chlorophyll a (chl-
122	$a L^{-1}$) using the non-acidification <i>in vitro</i> fluorometric technique (Welschmeyer,
123	1994). Seawater was filtered onto glass fiber filters (Whatman GF/F filters; 25-mm
124	diameter, 0.7 μ m nominal pore size) at low pressure (<70 kPa) and immediately
125	extracted in 90% acetone for approximately 24 h at -20°C. Chl-a concentrations were
126	determined with a Turner Designs 10AU fluorometer calibrated at the beginning of each
127	cruise with pure chl-a in 90% acetone and monitored for instrument drift during the
128	cruise using a solid secondary standard.
129	
130	2.5. Nutrients
131	Water samples for dissolved inorganic macronutrient analyses were collected at
132	a depth of 5 m using a 10 L Niskin bottle. Unfiltered samples were collected in pre-

133 cleaned polypropylene tubes and freshly analyzed at sea for nitrate plus nitrite $(NO_3^- + NO_3^-)$

134 NO_2^{-} ; hereafter referred to as nitrate), orthophosphate (PO₄³⁻), and silicate [Si(OH)₄]

135 with a Lachat QuikChem8000 Flow Injection Analysis system using standard

calorimetric techniques (Smith and Bogren, 2001; Knepel and Bogren, 2002; Wolters,
2002; respectively).

138

139 *2.6. Domoic acid*

140 Dissolved DA (dDA) and particulate DA (pDA) concentrations were measured 141 on sample filtrates and filters, respectively (Millipore Corp. mixed cellulose ester filters; 142 0.45 µm) using the direct competitive enzyme linked immunoassay (cELISA) Biosense 143 kits (Biosense Laboratories, Bergen, Norway), a modified version of the indirect 144 cELISA method as described in Garthwaite et al. (1998). Samples were analyzed in 145 duplicate, and the occasional poor replicates reanalyzed. The limit of detection for seawater samples was 6.8 ng L^{-1} , and the limit of quantification was 13.9 ng L^{-1} . 146 147 Cellular DA concentrations were estimated by dividing pDA concentrations by the 148 corresponding Pseudo-nitzschia cell concentrations. These estimates were restricted to 149 samples containing a minimum of 50 cells, providing a 95% confidence interval of \pm 150 30% of the mean cell density (Lund et al. 1958). As the focus on this paper is on DA 151 release by *Pseudo-nitzschia*, particulate DA along the drifter path data are not shown 152 but are the focus of a complementary paper by Lessard et al., (unpublished data). 153

154 2.7. <u>Pseudo-nitzschia</u> cell counts and species identification

Samples for cell quantification were collected at a nominal 1 m depth along the
Drifter A track using 10 L Niskin bottles. Total *Pseudo-nitzschia* cells were quantified
from whole water samples preserved with buffered formalin (<1% final concentration).
Cells were enumerated with a Palmer-Maloney counting chamber using a Zeiss

159 Axiovert 135 inverted light microscope. Samples (50 mL) were settled when necessary

160 for at least 24 h and counted at 200 x (total) magnification. Surface phytoplankton 161 samples were collected for *Pseudo-nitzschia* species identification at each station using 162 a 20-µm mesh phytoplankton net. P. cuspidata were positively identified using 163 transmission electron microscopy (Lundholm et al., 2003). 164 Eutreptiella cell counts. Sub-samples for cell quantification were collected from 165 the *Pseudo-nitzschia* sample bottles. Unpreserved, unstained *Eutreptiella* cells were 166 quantified immediately using a Becton Dickenson FACSCalibur flow cytometer, 167 equipped with a 15-mW laser exciting at 488 nm. Samples were run at a flow rate of 60 168 μ L min⁻¹ and the cells discriminated using particle size and chlorophyll content.

169

170 2.8. Sinking Rates

171 Samples for determination of sinking rates were collected from the surface 172 (approximately, 1 m depth) using 10 L Niskin bottles. A modified SetCol (settling 173 column) of Bienfang (1981) was used to estimate the sinking and floating rates of 174 phytoplankton (Beall, 2009). Triplicate ca. 500 mL samples of the natural assemblage 175 were homogenized and decanted into glass columns (2.6 cm diam., 50 cm tall). After a 176 3-h incubation, top, middle and bottom fractions were removed from the column. The 177 volumes of the fractions were measured for each settling column. The rates estimated 178 by the SetCol protocol were not sensitive to the fraction volumes as long as the top and 179 bottom fractions range from 10% to 15% of the total volume in the column (Bienfang, 180 1981). The biomass in each fraction was determined by chlorophyll extraction using the 181 same protocol as outlined previously. The sinking (ψ) and floating rates (A) were 182 calculated by Equation 1.

A or
$$\psi = \frac{B_{frac(obs)} - B_{frac(pred)}}{B_{tot}} \left(\frac{h}{t}\right)$$
 (Eq. 1)

where $B_{\text{frac}(\text{obs})}$ is the observed biomass in the top (floating) or bottom (sinking) 185 fractions. B_{frac(pred)} is the predicted biomass of the fraction based on the sum of 186 biomass in the column multiplied by the fraction volume relative to the total volume, h 187 is the height of the column of water and t is the incubation time in hours.

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189 2.9. Grow-out Incubation Studies

190 Seawater was collected every 2 days during the drift from a 5-10 m depth using a trace metal (TM) clean, all-Teflon[®] 'fish' pumping system that gently samples 191 192 phytoplankton and microzooplankton (Wells et al., 2009). All experimental preparations 193 were performed in a fabricated, positive pressure, all-plastic TM clean room using TM 194 clean techniques. Seawater was filtered through a pre-rinsed 200-µm nylon mesh (Nitex[®]) to remove grazers and homogenized in a 50-L polypropylene carboy. Nutrient 195 196 analysis was conducted on the initial seawater collected to determine if macronutrient 197 additions were required and amended as necessary to achieve 15 μ M nitrate, 2 μ M 198 orthophosphate, and 15 µM silicate at the beginning of the experiment. Trace metals 199 had previously been removed from macronutrient stocks using a Chelex-100 ion 200 exchange resin (Price et al., 1988; 1989). The thoroughly mixed seawater was dispensed 201 into 250-mL clear, polycarbonate bottles before treatments were added. Treatments 202 included amending the waters with iron (FeCl₃ additions of 1 or 3 nM), copper (CuSO₄ 203 at 1 or 3 nM) or reducing the free iron through the addition of the chelator

desferrioximine (DFB) (Desferal® - as deferoxamine mesylate is N-[5-[3-[(5-204 205 aminopentyl)hydroxycarbamoyl]propionamido]pentyl]-3-[[5-(N-206 hydroxyacetamido)pentyl]carbamoyl]propionohydroxamic acid monomethanesulfonate) at 3 nM. The fully prepared bottles were placed in a clear Plexiglas[®] incubators. 207 208 Temperature was maintained at sea-surface temperature with flowing surface seawater 209 and the incident photosynthetic photon flux density (PPFD) reduced to an equivalent of ~50% of the maximum daytime irradiation using a combination of neutral density 210 screening and blue Plexiglas[®]. Bottles were retrieved after four days (Day 4) and 211 212 processed.

213

214 2.10. Photosynthetic Efficiency

215 Photosynthetic efficiencies were measured by comparing the photosynthesis vs. irradiance responses (P vs. E) on Days 0, 2, 4, and 6. Rates of photosynthesis derived 216 from the amount of NaH¹⁴CO₃ incorporated into the cells during short-term incubations 217 218 in a temperature-controlled photosynthetron under controlled light intensities. Five-mL 219 subsamples were dispensed into 20-mL clear glass scintillation vials and each vial was inoculated with 5 μ Ci of NaH¹⁴CO₃. The vials were then strategically placed in a cool-220 white (halogen) light-field that provides PPFDs from 0 to 1,200 μ mol photons m⁻² s⁻¹. 221 222 After incubation, subsamples were acidified with 0.5 mL of 10% HCl (v/v) to stop the 223 reaction and allowed to degas for *ca*. 24 h prior to the addition of 15 mL of scintillation fluid (EcoLumeTM, MP Biomedicals LLC). Samples were subsequently mixed by gentle 224 225 inversion and allowed to sit undisturbed in the dark until they were radio-assayed using liquid scintillation counting. All ¹⁴C uptake (photosynthesis) rates were corrected for 226

227	dark uptake using the formula of Parsons et al. (1984), P-E curves generated using a
228	non-linear, least-squares regression technique (KaleidaGraph [©] ; Synergy Software).
229	Rate estimates of photosynthesis, normalized to chlorophyll a were fitted to the 3-
230	parameter P-E model of Platt and Gallegos (1980).
231	
232 233	3. Results
234	During September of 2004, the Juan de Fuca eddy region experienced high
235	densities of Pseudo-nitzschia spp. and the marine euglenoid, Eutreptiella sp.
236	Lagrangian ARGOS-tracked drifters, deployed at the origin of the upwelled nutrients of
237	the eddy, enabled us to continually track and sample the Pseudo-nitzschia bloom for
238	seven days from its formation to demise. The most common species of Pseudo-
239	nitzschia present was P. cuspidata that co-dominated the phytoplankton assemblage
240	with the euglenoid, Eutreptiella sp.
241	This particular drifter was one of several deployed during September 2004 as
242	part of a continuing evaluation of the surface current transport in the Strait of Juan de
243	Fuca to Columbia River regions (McFadyen et al., 2005). Once deployed in the
244	upwelled waters adjacent to the Juan de Fuca upwelling initiation site, the drifter
245	proceeded to move northwest for three days then shifted south and circled towards the
246	coastal waters often associated with the along-shore transport of the Columbia River
247	(MacFadyen et al., 2005). Prior to exiting the eddy, the drifter movement slowed from
248	drifter Day 4 to drifter Day 6 (Fig. 1). The short retention of the drifter corresponds to
249	an approach to the coastal front, where the water mass had distinctly different
250	temperature, salinity and nutrients. After drifter Day 6, the drifter continued moving in

a rapid pace southeast towards shore and was retrieved approximately 30 nautical milesoffshore.

253 Biological and chemical measurements of the surface waters indicated that as 254 the drifter proceeded through time and space, the sinuous path between drifter Day 5 255 and 6 corresponded to definable differences in water masses (Fig. 2). Water 256 temperatures first declined and then increased, with the drifter experiencing moderately 257 cooler, higher salinity waters on drifter Day 5 (Fig. 2A, 2B). From Days 1-4, total 258 biomass (chl-*a*) increased then decreased, while nitrate and silicate concentrations 259 increased slightly and the total densities of *Pseudo-nitzschia* spp. increased, then 260 remained fairly constant (Fig. 3A-C). Inexplicably, the total abundance of Pseudo-261 nitzschia decreased 3-fold on Day 5 (Fig. 3C), without a corresponding decrease in 262 biomass or drawdown of either nitrate or silicate (Fig. 3B). During the early days of the 263 drifter-based observations, the loss of *Pseudo-nitzschia* cells from the surface waters 264 was minimal. The cell sinking rates and the aggregation rates of *Pseudo-nitzschia* cells 265 remained inconsequential, ensuring that the cells remained in the surface waters (Fig. 4) 266 until the drifter and associated waters were impacted by the putative internal wave 267 upwelling event on drifter Day 5 (details in Lessard et al., in prep.). 268 The most dramatic changes in the biology and chemistry occurred around drifter 269 Day 5 associated with the cooler, higher salinity subsurface water signal (Figs. 2-3; 270 drifter Days 6-8). After the exchange with the vertically mixed water mass there was a 271 concurrent increase in cell aggregation (Fig. 4), a rapid increase in Pseudo-nitzschia-272 specific sinking rates, leading to a reduction in *Pseudo-nitzschia* cell density (Fig. 3C), 273 and an unexpected but prominent increase in dDA concentrations (Figure 3D).

274 Coincidentally, there was also a rapid drawdown of nitrate, but not silicate – indicating

275 a stimulated nutrient-drawdown by community members that did not include Pseudo-276 nitzschia or competing diatoms. Overall, total biomass was not reduced in the drifter days succeeding the transfer through the front. The most logical beneficiary of the 277 278 change in water mass was the co-blooming euglenoid, Eutreptiella sp., that matched the 279 cell densities of Pseudo-nitzschia for the first four drifter days (Lessard et al., in prep.). 280 In addition to the measurement of *Pseudo-nitzschia* density, macronutrient and 281 DA concentrations, and the floating/sinking rates of *Pseudo-nitzschia*, the physiological 282 "health" of the natural population was assessed using waters collected on Days 0, 2, 4 283 and 6 along the drifter path (Fig. 1). These waters were incubated for four days in deck-284 board incubators after amendment with Fe or Cu – elements that we propose influence 285 the growth and DA-levels in *Pseudo-nitzschia*. The resulting photo-physiological 286 "health" was assessed using the relative performance of the photosystem under 287 increasing photosynthetic photon flux density (PPFD; Fig. 5). The photosynthetic rates, 288 normalized to the concentration of *Pseudo-nitzschia* cells, were compared using 289 ambient water supplemented with either 3 nM DFB or 3nM Cu + 3 nM DFB. 290 Respectively, the enrichments allowed us to consider if the ambient waters were Fe-291 limited, Cu-limited, could be Fe-deplete and could be a combination of Fe-deplete but requiring Cu. We report the photo-physiological "health" of the community using the 292 maximum rate of photosynthesis (P^b), estimated from the P vs. E curves for the natural 293 294 phytoplankton community under natural conditions and then compared to amended 295 conditions. For example, control (ambient) water collected on Day 0 (Fig. 5) achieved a maximum photosynthetic rate of *ca*. 4.5 μ g C (μ g chl *a*)⁻¹ h⁻¹. When these natural 296 297 waters were amended with 3 nM DFB, creating Fe-stressed cells, the maximum 298 photosynthetic rate decreased ~50%; whereas, when the Fe-stressed cells were also

supplemented with 3 nM Cu, the maximum photosynthetic rate was over 4-fold that ofthe iron-stressed cells, and over 2-fold the rate of non-augmented cells.

301 Following this strategy, when cells along the drifter path were assessed for their 302 'photosynthetic health' (via assessment of the photo-physiological state) the early cells, 303 cells collected directly from the newly upwelled eddy waters, had a photosynthetic max of 4 μ g C (μ g chl a)⁻¹ h⁻¹ (Fig. 6), but the maximum photosynthetic rate dropped 304 dramatically to less than $2 \mu g C (\mu g chl a) h^{-1}$ for drifter Days 2 and 4. Supplementing 305 306 these waters with 3 nM Fe did not replenish the community's physiological deficiency. 307 However, by depleting the waters of available iron through the addition of DFB and adding 3 nM Cu maximum photosynthetic rates of 8-11 μ g C (μ g chl a)⁻¹ h⁻¹ were 308 309 achieved.

310 In the second phase of the drifter path, when high salinity, low temperature 311 waters altered the character of the surface waters (drifter Days 6-7; Fig. 2), the 312 physiological response of the community changed dramatically. The community 313 achieved greater maximum photosynthetic rates, with values approaching 7 μ g C (μ g chl a)⁻¹ h⁻⁻¹ (Fig. 6). Supplementing the community with 3 nM Fe neither increased nor 314 315 decreased the photosynthetic capacity. However, the amendment of the Fe-enriched 316 sample with 3 nM Cu depressed the maximum photosynthetic capacity to $< 1 \mu g C (\mu g)$ chl a)⁻¹ h⁻¹ – a reduction of rates that indicate Cu-toxicity. 317

To assess the impact of Cu and Fe additions, or Fe removal by DFB enrichment, 1 or 3 nM of either Fe, Cu, DFB or DFB + Cu were added to the grow-out incubation bottles, and cell growth and toxin characteristics were assessed after 4 days of deckboard incubation. At all stations, the addition of Fe-stimulated biomass proportional to

322 the level of Fe added – indicative of some degree of Fe limitation. In contrast, the 323 addition of Cu reduced the achieved biomass. The increase in biomass was positively 324 related to the concentration of added Fe (Fig. 7A). There was a corresponding reduction 325 in achieved biomass under all other treatments for drifter Days 0, 2, and 4. The 326 community from drifter Day 6 showed no significant difference in growth compared 327 with the control (open bars) when provided Cu, DFB or DFB + Cu (Fig. 7B-D). 328 Considering that the treatments may not affect all genera in a similar fashion, the 329 concentrations of *Pseudo-nitzschia* were recorded over the 4-day incubation period. 330 Generally, Pseudo-nitzschia was stimulated by all treatments, in particular after drifter 331 Day 0 (Fig. 8). The most dramatic stimulation of *Pseudo-nitzschia* growth occurred on 332 drifter Day 4, particularly when DFB was added, alone or in combination with Cu 333 (Figure 8B, D), whereas the Pseudo-nitzschia population from drifter Day 6 responded 334 weakly to all treatments.



- **Figure 1**. Path of the drifter in the Juan de Fuca Eddy during ECOHAB-PNW-III
- 341 Cruise: **A.** An overlay of the drifter path over a SeaWIFS image of the Juan de Fuca
- eddy region image taken Day 3 (September 19, 2004). **B.** The track of drifter. The
- 343 drifter was placed in the central edge of the eddy on Day 0. Samples for the data
- 344 presented here were collected at the indicated locations as red circles (•) using either a
- 345 10 L Niskin bottles or using the trace metal clean 'fish' and pump sampler.

348 13.4 **-**32.36 _T 7.0 В ۱C А 6.6 Temperature (°C) Oxygen (mL L⁻¹) 32.26 13.0 Salinity 6.2 12.6 32.16 5.8 ó 0 C 2 4 Drifter Day 2 4 Drifter Day 2 Drifter Day 6 . 6 6 0 0 0 349 350



Figure 2. Physical and chemical parameters measured on Days 0, 2, 4, and 6,

353 corresponding to the dates of the grow-out experiments, along the drifter path in the

Juan de Fuca Eddy, September 2004. Discrete samples were collected using 10-L

355 Niskin bottles and processed on board. The dashed line at Day 5 signifies entry into the

356 second phase of the drifter path.



Figure 3. Development of the A) phytoplankton biomass (extracted chlorophyll *a*) (n =
2, range reported), B) nitrate and silicate concentrations, C) Surface *Pseudo-nitzschia*concentrations and D) Surface dissolved DA during the drifter path in the Juan de Fuca
Eddy in September 2004. Samples were collected using 10 L Niskin bottles. The
dashed lines signify entry into the second phase of the drifter path after Day 5.



Figure 4. Phytoplankton community sinking rates (○) and aggregation (floating; ●)
rates in the mixed layer (5 m depth) during the Juan de Fuca drifter experiment in
September 2004. Samples were collected at the 5-m using 10 L Niskin bottles. The
dashed lines signify entry into the second phase of the drifter path after Day 5.



Figure 5. Effects of 3 nM desferoxamine (DFB) and 3 nM Cu additions on the
photosynthetic capacity of natural phytoplankton communities as a function of
photosynthetic photon flux density (PPFD). All samples collected from drifter Day 0
using the trace metal clean fish. The control represents photosynthetic rates from unamended waters.



Figure 6. Effects of Fe and Cu additions on the maximum photosynthetic rates of cells
collected on Days 0, 2, 4, and 6 along the drifter path using the trace metal clean fish.
The combination of Fe + Cu addition enhanced the photosynthetic performance in the *Pseudo-nitzschia* dominated early stages of the drifter path, whereas, the phytoplankton
community from the sample stations after drifter Day 4 were highly sensitive to trace
metal additions. Natural samples are unamended waters. The dashed lines signify entry
into the second phase of the drifter path after Day 5.



Figure 7. Responses of phytoplankton biomass measured after 4-day deck-board 'growout' incubations of ambient water collected using the trace metal clean fish on Days 0,
2, 4, and 6 along the drifter path in the Juan de Fuca Eddy, September 2004. Duplicate
bottles were supplemented with macronutrients, then further supplemented with either
A) iron, B) desferoxamine (DFB), C) copper, and D) copper + DFB. Bars show the
range of responses (n=2).



Figure 8. The growth response of *Pseudo-nitzschia* after 4-day 'grow-out' incubations
of ambient water collected using the trace metal clean fish on Days 0, 2, 4, and 6 along
the drifter path in the Juan de Fuca Eddy, September 2004. Bottles were supplemented
with indicated macronutrients, and with either: A) iron, B) desferoxamine (DFB), C)
copper, and D) copper + DFB. Single samples were analyzed.





412 Figure 9. In situ concentrations of particulate DA (pDA) verses ambient concentrations



414 PNW-III Cruise in September 2004 (data from Trainer et al., 2009).

4. Discussion

417	The Juan de Fuca Eddy off Vancouver Island, British Columbia, Canada and the
418	Washington coast, USA was reported by Trainer et al. (2009a, b) to contain a large,
419	nearly monospecific diatom bloom of P. cuspidata in September 2004 that also
420	contained the euglenoid, Eutreptiella spp. Pseudo-nitzschia cuspidata reached cell
421	densities of ~ 6.1 x 10^6 cells L ⁻¹ and produced maximum particulate domoic acid
422	(pDA), dissolved domoic acid (dDA), and cellular domoic acid concentrations of 43
423	nmol L^{-1} , 4 nmol L^{-1} , and 63 pg cell ⁻¹ , respectively. The synoptic survey conducted
424	during this time revealed that 84% of the stations ($n = 598$) had detectable <i>Pseudo</i> -
425	nitzschia cells and 78% had detectable levels of pDA. Variable ratios of pDA:dDA in
426	the eddy region suggested that DA release was under cellular regulation by Pseudo-
427	nitzschia, however, there were no significant correlations between either pDA or
428	cellular DA and ambient concentrations of macronutrients, including silicate (Figure 9).
429	Even so, pDA in surface waters (1-5 m depth) was positively correlated with chl a and
430	negatively correlated with temperature ($p < 0.01$) when <i>Pseudo-nitzschia</i> was present.
431	These findings demonstrate that Si limitation is not a prerequisite or 'trigger' for
432	Pseudo-nitzschia toxicity, as is commonly stated (e.g., Du et al., 2016), and that the
433	mechanistic basis for DA synthesis is linked instead to other environmental or
434	nutritional factors. Similarly, there were no significant correlations between cellular
435	DA concentrations and planktonic bacteria or cyanobacteria abundances (Beall, 2009),
436	contrary to the purported links between bacteria activity and DA production by Pseudo-
437	nitzschia (cf., review by Lelong et al. 2014), although we cannot rule out that there may
438	have been significant changes in the composition of the bacterial community (e.g.,

439 Hattenrath-Lehmann and Gobler, 2017) or that bacteria comprising the biofilm of 440 natural eukaryotic cell walls may contribute to DA production. There was a correlation between limiting concentrations of Fe (*ca*. 0.1 nmol L^{-1}) and the greatest *Pseudo*-441 442 nitzschia abundances, as well as pDA and dDA concentrations (Trainer et al., 2009a, b). 443 The study here focused on a single drifter deployment during the synoptic 444 survey described by Trainer et al. (2009a) where the oceanographic and physiological 445 conditions of a surface population of *Pseudo-nitzschia* were tracked over time within 446 the Juan de Fuca Eddy system. The drifter faithfully followed a single patch of surface 447 water over six days before being ejected in a southeast direction across the perimeter of 448 the eddy core on Day 7, requiring 2 more days to be ejected completely from the eddy 449 into the coastal current (Lessard et al., in prep.). Internal wave forcing within this 450 dynamic system generated periodic infusions of colder, more saline, subsurface waters 451 into the surface patch (Fig. 2), which would have resupplied it with nutrients. Here, our 452 findings can be viewed in terms of a natural "semi-continuous batch" culture system, 453 where the phytoplankton community in the advected surface patch was supported by at 454 least two nutrient re-infusions over the 6-day circumnavigation of the eddy core (Fig. 455 3).

The periodic vertical infusion of subsurface water helps to explain the rather sluggish changes in dissolved macronutrient concentrations that accompanied the small but marked increases in total biomass and significant increases in *Pseudo-nitzschia* abundance (Fig. 3). There was no net consumption of nitrate, silicate, and phosphate (data not shown) over this period, and indeed nitrate concentrations actually increased as the bloom developed (Fig 3B). The combination of a persistently elevated *Pseudonitzschia* abundance (Fig 3C), stable photosynthetic rates (Fig. 6), and low sinking rates

463 (Fig. 4), all point to the maintenance of a healthy phytoplankton community over the
464 first 4 days. Indeed, the initially elevated rates of aggregation and lower cell abundance
465 on Day 0 suggest that the *Pseudo-nitzschia* community had re-emerged from less
466 favorable pre-drifter conditions, perhaps stimulated by vertically-advected infusion of
467 subsurface waters prior to sampling and Day 1 (Fig. 1).

468 Pseudo-nitzschia cell densities declined precipitously between Days 4 and 5 469 (Fig. 3c), signaling the collapse of the bloom. The onset of this collapse was not closely 470 related to declining macronutrient concentrations, which occurred after Day 5. Bloom 471 termination corresponded closely with the intensification of Pseudo-nitzschia-cell-472 specific sinking rates (Fig. 4) and the order of magnitude increase in dissolved DA 473 concentrations (ca. 2 to 20 nM dDA) in surface waters (Fig. 4D). Nonetheless, rapid 474 declines in *Pseudo-nitzschia* abundance after Day 4 had little effect on total 475 (chlorophyll) biomass (Fig 3A) due to the compensatory increase in the abundance of 476 the marine euglenoid, Eutreptiella (Lessard et al., in prep.). 477 The factor, or factors, causing the transition from a Pseudo-nitzschia- to 478 Eutreptiella-dominated community after Day 4 are not known. There is no indication 479 that increased grazing pressure was responsible (Lessard et al., in prep.). And the 480 collapse of the *Pseudo-nitzschia* bloom clearly was not triggered by limiting 481 concentrations of nitrate, silicate, or phosphate; the acute drop in nitrate concentrations 482 occurred between Days 5 and 6, after the decline in Pseudo-nitzschia began, and even 483 on Day 6, nitrate concentrations were still in excess of those required to support 484 maximum rates of uptake by cells of the P. pseudodelicatissima complex (Auro and 485 Cochlan, 2013). This is in contrast with prior, largely stable nutrient conditions,

486 suggesting that nutrient infusion from internal wave processes ceased, consistent with

487 the physical data (Fig. 2). Despite the rapid decrease in *Pseudo-nitzschia* abundance 488 there were only marginal if any change in total biomass (Fig. 3b), attributable to further 489 in-growth of *Eutreptiella* spp. (Lessard et al., in prep.). The enhanced growth of 490 *Eutreptiella* within the community coincides with a roughly 3-fold increase in 491 photosynthetic capacity of the transitioning community (Fig 6) and the relaxing of the 492 iron-limitation of the community. The absence of comparable silicate drawdown (Fig. 493 3b) is further evidence of the declining health of *Pseudo-nitzschia* during this transition. 494 If not decreased availability of macronutrients, what then could have initiated 495 the collapse of the Pseudo-nitzschia bloom? It is feasible that the transition was 496 associated with some degree of lateral advection of "patchy" surface waters, given the 497 dynamics of this system, however, the survey data showed reasonable spatial uniformity 498 in the composition of surface water phytoplankton community (Trainer et al., 2009a). 499 At best, any lateral advection would have accelerated the apparent community transition 500 that was occurring more broadly in the region. 501 The deck-board incubation experiments demonstrate that the community was Fe 502 stressed, with the Fe treatment generating markedly greater total biomass (Fig 7a) and 503 Pseudo-nitzschia abundance (Fig. 8a) over the course of the drifter study. The same was 504 true immediately outside the core perimeter of the Eddy on Day 6 (Fig. 7a). 505 Importantly Fe amendments did not enhance cell photophysiology (Fig. 6) however, 506 indicating that there was sufficient Fe availability in the surface waters to maintain the 507 existing community, but that more Fe was needed to enable further biomass 508 accumulation. Even so, the abnormally elevated dissolved DA concentrations (ca. 2-3

509 nM) are consistent with *Pseudo-nitzschia* experiencing significant Fe stress (Maldonado

510 et al., 2002; Wells et al., 2005).

511 Further evidence that there was insufficient biologically-available Fe during the 512 drifter bloom is the comparative ease that Fe stress could be induced by adding the 513 siderophore DFB. Additions of DFB to both laboratory cultures and natural 514 phytoplankton communities can decrease Fe availability (Hutchins et al., 1999; Wells, 515 1999; Wells and Trick, 2004, Shaked and Lis, 2012), and the same decreasing trend in 516 achieved total biomass with progressively greater siderophore concentrations was seen 517 here (Fig. 7b). The increased Fe stress is seen in the lowered community photosynthetic 518 capacity in the DFB- amended incubations (Fig. 5). Although dissolved Fe 519 concentrations were not measured during the cruise, previous experience in the region 520 suggested that 1-3 nM DFB additions would complex 'some-to-much' of the dissolved 521 Fe pool, through ligand exchange with the natural organic ligands (Rue and Bruland, 522 2001). But trace metal determinations during three subsequent cruises to this region 523 showed Fe concentrations of ca. 2-5 nM Fe (Roy, 2009), so it is reasonable to expect 524 that the DFB amendments would have been sufficient to complex much but perhaps not 525 all of the ambient dissolved Fe. This perspective is in agreement with the somewhat 526 muted impact of DFB on total biomass accumulation (Fig. 7B). In summary, the 527 phytoplankton community was Fe-stressed at the start and over the 6-day drifter 528 experiment, with biomass increasing upon Fe addition, and decreasing when a portion 529 of the ambient pool of Fe was made less available by complexation with DFB. 530 DFB amendments led to a decreased abundance of Pseudo-nitzschia in the deck-531 board incubations during the early phases of the bloom (Days 2 & 4), but this effect was 532 vividly reversed on Day 6, when Pseudo-nitzschia abundance increased sharply in the 533 DFB treatment (Fig. 8a, b). This reversal coincided with onset of the bloom decline in 534 surface waters (Fig. 3C), and the order of magnitude increase in dissolved DA

535	concentrations (Fig. 3D). So, when the deck-board incubation was started on Day 6, DA
536	concentrations were > 5-fold in excess of the added DFB concentration. Although DA
537	complexes Fe in seawater (Rue and Bruland, 2001), it still would not have competed
538	effectively with DFB for Fe given the massive difference in their conditional stability
539	constants in seawater $(log K_{FeDA,Fe(III)}^{cond} = 10^{8.7 \pm 0.5} \text{ M}^{-1} \text{ vs. } log K_{FeDFB,Fe(III)}^{cond} = 10^{16.5},$
540	Rue and Bruland, 1995, 2001). It is likely instead that the increased dDA
541	concentrations signal physiological impacts on Pseudo-nitzschia associated with the
542	increased Fe stress caused by DFB. A similar response has been shown in culture when
543	Pseudo-nitzschia spp. are placed under Fe limiting conditions (Wells et al., 2005).
544	P. multiseries and P. australis can fulfil their Fe growth requirements for rapid
545	exponential growth when non-complexed dissolved Fe concentrations (Fe') are greater
546	than ~ 25 pM (Maldonado et al., 2002), and it is reasonable to anticipate that P .
547	cuspidata would not have substantially different requirements. The bulk dissolved Fe
548	concentrations in these waters would be in far excess of this oceanic level (Roy, 2009),
549	meaning that the vast bulk of dissolved Fe in the drifting patch existed in organically
550	complexed forms that were not easily available to P. cuspidata (Trick et al., 2004).
551	Although Pseudo-nitzschia sp. appear to be less efficient at Fe uptake than other
552	diatoms under Fe-replete conditions, experiments demonstrate their superior ability to
553	adapt under Fe deplete conditions (Maldonado et al., 2002; Wells et al., 2005), and that
554	the production and release of DA facilitates Fe acquisition and alleviating Fe stress.
555	This enhanced Fe acquisition appears to result from Pseudo-nitzschia being able to
556	directly extract Fe from strong Fe complexes, as long as Cu is readily available
557	(Maldonado et al., 2002 Wells et al., 2005). Other diatoms also show a Cu requirement

558	for enhanced Fe uptake under Fe stressed conditions, along with reduced Fe
559	requirements through substitution for Fe-containing enzymes in the photosynthetic
560	transport chain (Grotz and Guerinot, 2006; Nouet et al., 2011). Here, Cu amendment
561	had little to no consistent effect on total biomass, yielding only slight decreases at the
562	highest Cu addition (3 nM) on Days 2 and 4 (Fig. 7C). However, adding Cu sharply
563	stimulated Pseudo-nitzschia growth within the community (Fig 8C) when Pseudo-
564	nitzschia abundance was high (Fig 3C). The same result has been observed in cultures
565	under Fe stress conditions (Maldonado et al., 2002; Wells et al., 2005).
566	The addition of Cu with DFB enhanced Pseudo-nitzschia growth during the
567	earlier phase of the bloom by 30-45% over that observed in the DFB only treatment
568	(Fig. 8C, D). This effect was even more clearly shown in the photo-physiological
569	measurements of the <i>in-situ</i> assemblage, where addition of Cu with DFB reversed the
570	negative effect of DFB alone (Fig. 5); indeed, the combination of DFB and Cu yielded a
571	higher photosynthetic capacity than the control. However, these positive effects were
572	reversed in the incubation initiated on Day 5, when the collapse of the bloom had
573	begun, where the combination of decreased Fe availability and increased Cu additions
574	led to markedly lower Pseudo-nitzschia growth. It may be that the decreases in health
575	of the Pseudo-nitzschia population, as shown by the increased aggregation, sinking
576	rates, and photo-physiology, were sufficiently massive to prevent recovery.
577	The substantial increase in dDA at the end of the bloom would not have
578	substantially affected dissolved Fe speciation. Rue and Bruland (2001) calculate that
579	dDA concentrations of 100 nM would lead to DA complexation of ca. 25% of the
580	dissolved Fe pool, so by extrapolation only ca. 5% of Fe would have existed as a DA
581	complex in the drifter surface waters. However, the DA release would have affected

582 dissolved Cu speciation to a much greater extent, reducing free cupric ion 583 concentrations by a factor of 2-3 during Days 0-4, and by more than a 100-fold after 584 Day 5 (Rue and Bruland, 2001). Given the combined observations showing targeted 585 release of DA by Fe-starved Pseudo-nitzschia, it's strong effect on Cu complexation, 586 the positive Cu effect on Pseudo-nitzschia photo-physiology (Fig. 5), and the necessity 587 of Cu for enhancing growth of Fe-stressed Pseudo-nitzschia in both the laboratory 588 (Maldonado et al., 2002; Wells et al., 2005) and field conditions (Wells et al., 2005; Fig. 589 8C), we believe that DA production affects the ecophysiology and success of Pseudo-590 nitzschia in coastal waters through the alleviation of Fe limitation via enhanced Cu 591 acquisition. 592 The sharp increase in dDA concentrations coincided with the transition from a 593 Pseudo-nitzschia-dominated to a Eutreptiella-dominated community. Pseudo-nitzschia 594 spp. appear to be somewhat more sensitive to Cu toxicity than other diatoms 595 (Maldonado et al., 2002), and one might hypothesize that the purpose of DA release by 596 the cell is to diminish Cu toxicity, were it not for the observations that Cu is essential 597 for inducing the high affinity Fe uptake system in *Pseudo-nitzschia* (Wells et al., 2005). 598 Although we have no direct measure of the sensitivity of the co-occurring *Eutreptiella* 599 spp. to Cu, other euglenoids show significant sensitivity (Netto et al., 2012). It seems 600 possible then that decreases in cupric ion concentrations due to the large release of 601 dissolved DA during the collapse of the Pseudo-nitzschia bloom may have enhanced the 602 growth success of Eutreptiella. 603 Nitrogen measurements on the cruise were limited to oxidized N forms, so we

are unable to assess whether DA production was related to changes in the supply of
 reduced nitrogen forms, even though we now know that the form of nitrogen is critical

to DA production (Howard, et al., 2007; Radan and Cochlan, 2018).

607 **5. Conclusions**

608 The rapid response to additions of Cu and Fe - leading to increased biomass -609 changes that are catalysed with the release of dDA, illustrate why simple correlations 610 between dDA and environmental conditions, such as with macronutrient concentrations, 611 have not revealed a strong association. Our multi-day investigation of bloom 612 progression illustrates that the rapid release of dDA is highest at the intersection of three 613 critical conditions (sufficient macronutrients, low Fe and low Cu). Our corresponding 614 survey approach never revealed the intensive release of dDA under these conditions. 615 The lack of reveal is due potentially to the rapid response of the cells to altered 616 environmental condition, and the ephemeral nature of the three conditions in the waters 617 of the PNW. Typical static cruise sampling protocols where the time (distance) 618 between sampling locations are long lack to precision to capture the toxin signal. This 619 work adds supporting evidence to the important role of trace metals, not macronutrient 620 limitation (most notably silicate) in forecasting the success and demise of toxigenic 621 Pseudo-nitzschia in the PNW. Although the toxic threat of DA to coastal ecosystems 622 and the health of marine mammals, birds and humans, is directly linked to the 623 concentration of particulate DA (pDA). The present study clearly underscores the 624 physiological importance of dissolved DA in the development of such toxic diatom 625 blooms. It is not just a wicked problem, but a wickedly transient problem. 626

627

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