

## Biodiversity and ecosystem functioning in stressed environments: primary producers and consumers at the basis of marine food webs

Christoph Mensens

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# Biodiversity and ecosystem functioning in stressed environments: primary producers and consumers at the basis of marine food webs

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## Table of contents

Summary		1
Samenvatting		5
Chapter 1	General Introduction	10
Chapter 2	Stressor-induced biodiversity gradients: revisiting biodiversity – ecosystem functioning relationships	31
Chapter 3	Different response-effect trait relationships underlie contrasting numerical and functional responses to two chemical stressors	45
Chapter 4	Facilitation and tolerance explain the diversity effect on functioning in stressed diatom communities	62
Chapter 5	Selective and context-dependent effects drive chemical stressor impacts across trophic levels at the basis of marine food webs	78
Chapter 6	General Discussion	97
Addenda	Addendum I – Supporting Material Chapter 2	115
	Addendum II – Supporting Material Chapter 3	123
	Addendum III – Supporting Material Chapter 4	141
	Addendum IV – Supporting Material Chapter 5	147
	Addendum V – Acute toxicity of copper to harpacticoid copepods	152
References		154

#### 1 Summary

Biodiversity is increasingly altered by human activities, and there is a growing concern to what
extent biodiversity loss will affect ecological processes that are essential to ecosystem functioning
and ultimately to human well-being. These concerns have made the relationship between
biodiversity and ecosystem functioning (diversity-functioning hereafter) a core research area
in ecology, and two decades of intensive research have provided compelling evidence for a link
between biodiversity and ecosystem functioning.

8 Anthropogenic stressors such as an increasing number of chemical contaminants are the driving 9 force behind the ongoing biodiversity loss. Whilst diversity-functioning research is often framed 10 within the context of anthropogenic change, anthropogenic stressors are however rarely included 11 into the design of diversity-functioning studies. Similarly, the majority of diversity-functioning 12 experiments have focussed on single trophic levels, whereas the impact of anthropogenic 13 stressors across trophic levels has remained underrepresented in diversity-functioning research.

14 This PhD used marine primary producers (benthic diatoms) and consumers (harpacticoid 15 **copepods**) and two chemical stressors (atrazine and copper) to explore four central questions 16 which are still largely debated or unresolved in current diversity-functioning research: (i) Does 17 the classic design in diversity-functioning research, which consists of random species loss, predict 18 the diversity gradients and diversity-functioning relations induced by anthropogenic stress? (ii) 19 Can biodiversity loss under stress, and its impact on ecosystem functioning, be predicted from the 20 species' traits? (iii) Is the biodiversity effect on functioning in stressed communities determined 21 by the dominant species (dominance effect) or a better performance of species in community 22 (complementarity effect), and which are the mechanisms driving both effects? (iv) Are stress 23 effects on the energy content of primary producers (diet quality) transmitted to the consumer 24 level? The main findings regarding these questions are presented in Chapters 2 to 5, whilst 25 Chapter 6 proposes an integrative synthesis of the three key effects through which anthropogenic 26 stress drives the diversity-functioning relation.

27 The majority of diversity-functioning experiments to date used broad gradients of species 28 richness, and tested the effect of biodiversity loss on ecosystem functioning by removing species 29 in a random way. It remains unsure however whether this classic protocol is representative of the 30 biodiversity loss and diversity-functioning relations as induced by anthropogenic stress. The first 31 objective of this work was therefore to contrast the biodiversity gradients and diversity-32 functioning relations in a benthic diatom community as obtained with the classic random design 33 with those induced by atrazine (Chapter 2). Biodiversity was quantified as species richness and 34 evenness, and functioning as the diatom's contribution to primary production, biomass 35 production, energy content and sediment stabilization. Biodiversity gradients induced by atrazine 36 exposure were narrower than predicted by the classic random approach, since the herbicide 37 altered evenness but did not cause any species loss. Diversity- functioning relations for sediment 38 stabilization and energy content were steeper in stressed than in randomly assembled 39 communities. This disproportionately large decrease of functioning with stressor-induced 40 biodiversity loss was related to selective atrazine effects on the species contributing most to 41 energy content and sediment stabilization (*Nitzschia* sp.), which was also most sensitive to stress. 42 The common approach in diversity-functioning research thus overestimated biodiversity loss 43 induced by chemical stressors, but underestimated the associated loss of function due to selective 44 stress effects targeting the species contributing most to functioning.

45 Given the plethora of anthropogenic stressors, it is logistically impossible to predict selective 46 stress effects by experimentally testing the tolerance of species to each of these stressors. Chapter 47 **3** therefore tested if response traits and effect traits could respectively predict the abundance 48 (numerical response) and functioning (functional response; potential contribution to primary 49 production, sediment stabilization and energy content) of 18 marine benthic diatom strains to 50 copper and atrazine. Furthermore, it was tested whether the species' numerical stress response 51 was related to their functional contribution and to their functional stress response, i.e. if atrazine 52 and copper targeted species of little or high importance for ecosystem functioning, and if tolerant 53 species could maintain their functional contribution under stress. The numerical and functional 54 response of diatoms to copper was predicted by the same set of intercorrelated morphological 55 traits (cell size, surface to volume ratio, cell length), with large cells being more resistant to the 56 metal. Under copper, the diatoms' numerical response stress response was positively related to 57 their functional contribution and functional stress response, meaning that copper-tolerant 58 species contributed most to functioning and performed best at maintaining their functioning. 59 Under atrazine, the capacity for mixotrophic growth predicted the numerical but not the 60 functional stress response of diatoms, and the diatoms' numerical response to atrazine was not 61 related to their functional contribution and functional stress response. Diatoms capable of 62 growing on organic carbonic sources could thus maintain their cell numbers, but contributed little 63 to functioning under herbicide stress. Overall, Chapter 3 indicated that if numerical and functional 64 stress response are predicted by the same set of response and effect traits, numerically tolerant 65 species might be able to maintain ecosystem functioning under stress. If response and effect traits 66 do not correspond, stress might cause a disproportionate loss of functioning.

Dominance (dominance by species with high or low functional contribution) and
complementarity (species contribute more to functioning in community than in monoculture) are
the two key effects driving the biodiversity effect on ecosystem functioning, but there is little

70 evidence on the effects of anthropogenic stress on dominance and complementarity. 71 Complementarity effects arise from ecological mechanisms such as niche partitioning or 72 facilitative interactions between species. Direct functional and ecophysiological tests of these 73 mechanisms are however scarce and there is no evidence attributing changes of complementarity 74 under stress to changing facilitative mechanisms. Chapter 4 first examined whether the 75 biodiversity effect on diatom biomass production under atrazine and copper stress was driven by 76 dominance or complementarity effects. Next, it was tested if the release of extracellular polymers 77 as a facilitative mechanism could explain changes in complementarity under stress. Last, it was 78 analysed if complementarity was two-way or one-way, i.e. if complementarity was equally 79 distributed among species, or favoured species depending on their properties. The biodiversity 80 effect on diatom biomass production depended on two-way complementarity, which increased 81 under stress. Under atrazine and copper stress, diatom communities increased their production 82 of extracellular polymers which, in part, predicted increases in complementarity. Diatom species 83 benefited from complementarity depending on their properties, with complementarity under 84 atrazine and copper respectively favouring the growth of mixotrophic (C. fusiformis, Nitzschia sp., 85 N. acicularis) and copper-sensitive species (A. lineolata, N. digitoradiata, Gyrosigma sp.). These 86 copper-sensitive species were however characterised by a low biomass production, causing a 87 negative one-way complementarity, which largely offset positive two-way complementarity 88 effects and limited the biodiversity effect on diatom biomass production under metal stress. 89 Chapter 4 provided first ecophysiological evidence for facilitation as driver of complementarity 90 under stress, by identifying extracellular polymers as a 'sleeping' facilitative mechanism, which 91 when stimulated under stress predicted the diversity effect on biomass production. This 92 facilitative mechanism was however not necessarily beneficial for ecosystem functioning when it 93 caused a negative one-way complementarity which benefited species with a low functional 94 contribution.

95 There is little evidence whether stressors drive functioning through selective effects (changes in 96 community structure) or context-dependent effects (changes in the species' functional 97 contribution), and if both types of stress knock on across trophic levels. Moreover, trophic 98 approaches to date have essentially focussed on diet quantity, i.e. primary producer biomass, 99 whilst diet quality (producer energy content) has not been considered when examining food web 100 functioning under stress. Chapter 5 measured if atrazine and copper affected functioning in a 101 diatom community (diet quantity, sediment stabilization, diet quality) through selective stress 102 effects (by selectively targeting the species contributing most to functioning) or context-103 dependent effects (by changing the species' functional contribution). Concomitant knock-on 104 effects of selective and context-dependent stress effects on the next trophic level were quantified 105 by testing the response of the diatoms' main grazer (the harpacticoid copepod *M. littorale*) to

106 changes in diet quality. Diatom diet quantity was reduced by copper stress but not by low atrazine 107 levels due to the presence of an atrazine-tolerant, mixotrophic species (C. closterium). The 108 diatoms' contribution to sediment stabilization was stimulated by context-dependent effects of 109 both stressors. At low atrazine levels, selective changes in community structure involving 110 dominance by the atrazine-tolerant but energy-poor species *C. closterium*, reduced diet quality by 111 more than half. Context-dependent stress effects only reduced diet quality at high atrazine and 112 copper levels. Furthermore, selective and context-dependent stress effects on diet quality affected 113 the energy transfer to the next trophic level, with *M. littorale* losing half of its energy content when 114 feeding on diatoms grown under atrazine and high copper stress. Chapter 5 identified selective 115 stress effects, causing shifts in community structure towards dominance by species with low 116 functional contribution, as a more potent threat for ecosystem functioning than any direct stress 117 effects on the species' functional contribution. The energy content of copepods depended on that 118 of their diatom diet, highlighting the relevance of diet quality as a key driver of energy transfer at 119 the primary producer-consumer interface.

120 **Chapter 6** integrates the three main effects influencing ecosystem functioning under stress: the 121 functional impact of selective stress, targeting the functionally most important species, increases 122 with differences in the species' functional contribution, and can be predicted from the species' 123 response and effect traits. Physiological stress, by directly reducing the species' functional 124 contribution, drives ecosystem functioning through context-dependent effects at high stress 125 levels. Increases in complementarity, driven by the activation of facilitative mechanisms under 126 stress, can enable primary producers to maintain their functional contribution and limit losses of 127 ecosystem functioning. By highlighting the three key effects driving the diversity-functioning 128 relation under stress, this work proposes an conclusive framework for a better integration of 129 anthropogenic stressors within diversity-functioning research.

#### 131 Samenvatting

132 Biodiversiteit wordt in toenemende mate veranderd door menselijke activiteiten, en er is een 133 groeiende bezorgdheid in hoeverre het verlies aan biodiversiteit invloed zal hebben op 134 ecologische processen die essentieel zijn voor het functioneren van ecosystemen en uiteindelijk 135 voor het menselijk welzijn. Deze bekommering zorgde ervoor dat de relatie tussen biodiversiteit 136 en het functioneren van ecosystemen (diversiteit - ecosysteem functionering) een 137 kernonderzoeksgebied in de ecologie werd, en twee decennia van intensief onderzoek hebben 138 overtuigend bewijs geleverd voor een link tussen biodiversiteit en het functioneren van 139 ecosystemen.

140 Antropogene stressoren zoals een toenemend aantal chemische verontreinigingen zijn de 141 drijvende kracht achter het voortdurende verlies aan biodiversiteit. Terwijl onderzoek naar 142 diversiteit - ecosysteem functionering vaak gekaderd wordt in de context van antropogene 143 veranderingen, worden de antropogene stressoren waardoor het voortdurende verlies aan 144 biodiversiteit echter zelden opgenomen in het ontwerp van diversiteit-functioneringsstudies. De 145 meeste diversiteit-functioneringsexperimenten richten zich op één trofische niveau, terwijl de 146 invloed van antropogene druk over trofische niveaus heen ondervertegenwoordigd is in 147 diversiteit-functioneringsonderzoek.

148 Dit doctoraat gebruikt mariene primaire producenten (benthische diatomeeën) en grazers 149 (harpacticoide copepoden) en twee chemische stressoren (atrazine en koper) om vier centrale 150 vragen die nog steeds grotendeels worden betwist of onopgelost zijn in het huidige onderzoek te 151 toetsen: (i) voorspelt het klassieke ontwerp in diversiteit-functioneringsonderzoek, dat bestaat 152 uit een willekeurig soortenverlies, de diversiteitsgradiënten en diversiteit-functionering relaties 153 veroorzaakt door antropogene stress? (ii) kan verlies van biodiversiteit onder stress en zijn 154 impact ervan op het functioneren van ecosystemen, worden voorspeld op basis van kenmerken 155 van de soort ? (iii) is het biodiversiteitseffect op het functioneren van gemeenschappen onder 156 stress bepaald door de dominante soort (dominantie-effect) of een betere prestatie van soorten 157 in de gemeenschap (complementariteitseffect), en welke zijn de mechanismen achter beide 158 effecten? (iv) worden stresseffecten op de energie-inhoud van primaire producenten 159 (dieetkwaliteit) doorgegeven naar het niveau van de consument? De belangrijkste bevindingen 160 ten aanzien van deze vragen worden gepresenteerd in de hoofdstukken 2 tot 5, terwijl hoofdstuk 161 6 een integratieve synthese geeft van de drie belangrijkste effecten waardoor antropogene stress 162 de diversiteit-functioneringsrelatie drijft.

Het merendeel van de diversiteit-functioneringsexperimenten gebruikte tot op heden een ruime
gradiënt van de soortenrijkdom, en testte het effect van het verlies aan biodiversiteit op het

165 functioneren van ecosystemen door het verwijderen van soorten op een willekeurige manier. Het 166 blijft echter onzeker of dit klassieke protocol representatief is voor het verlies aan biodiversiteit 167 en de diversiteit-functioning relaties veroorzaakt door antropogene stress. De eerste doelstelling 168 van dit onderzoek was dan ook om de biodiversiteitsgradiënten en diversiteit-169 functioneringsrelaties te contrasteren in een benthische diatomeeëngemeenschap zoals 170 verkregen met de klassieke willekeurig ontwerp versus een gemeenschap geïnduceerd door 171 atrazine (Hoofdstuk 2). Biodiversiteit werd gekwantificeerd als soortenrijkdom en 'evenness' 172 (spreiding van de individuen over de soorten), en functionering in termen van de bijdrage van de 173 diatomeeën aan de primaire productie, de productie van biomassa, energie-inhoud en 174 sedimentstabilisatie. Biodiversiteitsgradiënten die veroorzaakt werden door 175 atrazineblootstelling waren kleiner dan voorspeld door de klassieke willekeurige aanpak, omdat 176 het herbicide de spreiding van de individuen over de soorten ('evenness') wijzigde maar geen 177 verlies aan soorten veroorzaakte. Diversiteit-functioneringsrelaties voor sedimentstabilisatie en 178 energie-inhoud waren steiler in gestresseerde dan in willekeurig samengestelde gemeenschappen. Deze onevenredig grote vermindering van functioneren met stress-179 180 geïnduceerd verlies van biodiversiteit was gelinkt aan selectieve atrazine effecten op de soort die 181 het meest bijdroeg tot de energie-inhoud en sedimentstabilisatie (*Nitzschia* sp.). Dit was tevens 182 de meest stressgevoelige soort. De algemene aanpak in diversiteit-functioneringsonderzoek 183 overschat dus het verlies aan biodiversiteit veroorzaakt door chemische stressoren, maar 184 onderschat het daarmee gepaard gaande verlies van functie als gevolg van selectieve 185 stresseffecten gericht op de soorten die het meest bijdragen tot het functioneren.

186 Gezien de overvloed van antropogene stressoren is het logistiek onmogelijk om selectieve 187 stresseffecten te voorspellen door de tolerantie van soorten experimenteel te testen voor elk van 188 deze stressoren. Hoofdstuk 3 testte daarom of de responseigenschappen en effecteigenschappen 189 respectievelijk de abundantie (numerieke respons) en het functioneren (functionele respons; 190 potentiële bijdrage aan de primaire productie, sedimentstabilisatie en energie-inhoud) kunnen 191 voorspellen van 18 mariene benthische diatomeeënstammen tegenover koper en atrazine. Verder 192 werd onderzocht of numerieke stressreactie van soorten gerelateerd was aan hun functionele 193 bijdrage en hun functionele stressrespons. De numerieke en functionele respons van diatomeeën 194 ten opzichte van koper werd voorspeld door dezelfde set gecorreleerde morfologische 195 kenmerken (celgrootte, oppervlakte-volumeverhouding, cellengte), met grote cellen die beter 196 bestand zijn tegen koper. In de koperbehandelingen was de numerieke stressrespons van de 197 diatomeeën positief gerelateerd aan hun functionele bijdrage en functionele stressrespons, wat 198 betekent dat de koper-tolerante soorten het meest bijgedroegen tot het functioneren en het best 199 presteerden bij het handhaven van hun functie. Bij de atrazinebehandelingen voorspelde de 200 mixotrofe groei de numerieke maar niet de functionele stressreactie van diatomeeën en de 201 numerieke respons van de diatomeeën ten opzichte van atrazine was niet gerelateerd met hun 202 functionele bijdrage en functionele stressreactie. Diatomeeën die kunnen groeien op organische 203 koolzuurbronnen kunnen dus celdensiteiten behouden, maar dragen weinig bij tot het 204 functioneren onder herbicidestress. Over het algemeen, onderstreept Hoofdstuk 3 dat als 205 numerieke en functionele reactie op stress worden voorspeld door dezelfde set van de respons-206 en effecteigenschappen, numeriek tolerante soorten mogelijks het ecosysteem functioneren 207 kunnen behouden onder stress. Als respons- en effecteigenschappen niet overeenstemmen dan 208 kan stress een onevenredig verlies van functioneren veroorzaken.

209 Dominantie (dominantie van soorten met een hoge of lage functionele bijdrage) en 210 complementariteit (soorten die meer bijdragen aan het functioneren in de gemeenschap dan in 211 monocultuur) zijn de twee belangrijkste effecten die het effect van biodiversiteit op 212 ecosysteemfunctionering sturen maar er is weinig bewijs voor de effecten van antropogene druk 213 op dominantie en complementariteit. Complementariteitseffecten vloeien voort uit de ecologische 214 mechanismen zoals nichepartitionering of faciliterende interacties tussen soorten. Directe 215 functionele en ecofysiologische testen van deze mechanismen zijn echter schaars en er is geen 216 bewijs die wijzigingen in complementariteit onder stress kan toewijzen aan veranderende 217 faciliterende mechanismen. Hoofdstuk 4 onderzocht eerst of het biodiversiteitseffect op de 218 productie van biomassa door diatomeeën onder atrazine- en koperstress werd gedreven door 219 dominantie of complementariteitseffecten. Vervolgens werd onderzocht of de afgifte van 220 extracellulaire polymeren als faciliterende mechanisme een verklaring kan zijn voor 221 veranderingen in complementariteit onder stress. Tenslotte werd geanalyseerd of de 222 complementariteit in twee richtingen of één richting voorkwam, met andere woorden of de 223 complementariteit evenredig verdeeld werd tussen soorten, of of soorten bevoordeeld werden op 224 basis van hun eigenschappen. Het biodiversiteitseffect op de productie van biomassa door 225 diatomeeën was afhankelijk van twee-richtingen complementariteit, die toenam onder stress. 226 Onder atrazine en koperstress verhoogden de diatomeeëngemeenschappen hun productie van 227 extracellulaire polymeren wat deels de toename in complementariteit voorspelde. 228 Diatomeeënsoorten profiteerden van complementariteit afhankelijk van hun eigenschappen, 229 waarbij complementariteit bij behandeling met atrazine en koper de groei van respectievelijk 230 mixotrofe (C. fusiformis, Nitzschia sp., N. acicularis) en kopergevoelige soorten (A. lineolata, N. 231 digitoradiata, Gyrosigma sp.) promootte. Deze kopergevoelige soorten werden echter gekenmerkt 232 door lage biomassa-opbrengsten, waardoor een negatieve éénrichtingscomplementariteit, die 233 grotendeels positieve tweerichtingscomplementariteit effecten teniet deed en beperkte het 234 biodiversiteitseffect op de productie van diatomeeënbiomassa bij stress door metalen. Hoofdstuk 235 4 leverde het eerste ecofysiologisch bewijs voor facilitatie als drijfveer van complementariteit 236 onder stress, door het aanduiden van extracellulaire polymeren als een 'slapend' faciliterend 237 mechanisme, dat wanneer gestimuleerd onder stress het diversiteitseffect op de 238 biomassaproductie voorspelde. Deze faciliterende mechanisme was echter niet per se gunstig 239 voor het functioneren van het ecosysteem indien het een negatieve 240 éénrichtingscomplementariteit veroorzaakte die soorten bevoordeligde met een lage functionele 241 bijdrage.

242 Er is weinig bewijs dat stressoren functionering sturen door middel van selectieve effecten 243 (veranderingen in de gemeenschapsstructuur) of context-afhankelijke effecten (veranderingen in 244 functionele bijdrage van de soorten), en of beide types van stress een impact hebben over 245 trofische niveaus heen. Bovendien hebben trofische benaderingen tot op heden zich voornamelijk 246 gericht op dieethoeveelheid, dat wil zeggen de biomassa van de primaire producent, terwijl de 247 kwaliteit van het dieet (energie-inhoud van de producent) niet beschouwd werd bij het onderzoek 248 van het functioneren van een voedselweb onder stress. In Hoofdstuk 5 werd onderzocht of 249 atrazine en koper een effect hadden op het functioneren van een diatomeeëngemeenschap 250 (dieethoeveelheid, sedimentstabilisatie, dieetkwaliteit) door middel van selectieve stresseffecten 251 (door selectieve impact op de soorten die het meest bijdragen tot het functionering) of context-252 afhankelijke effecten (door het veranderen van de functionele bijdrage van de soorten). 253 Gelijktijdig knock-on effecten van selectieve en context-afhankelijke stress effecten op de 254 volgende trofische niveau werden gekwantificeerd door het testen van de respons van de 255 belangrijkste grazers op de diatomeeën (de harpacticoide copepode *M. littorale*) ten opzichte van 256 veranderingen in de dieetkwaliteit. De hoeveelheid van het diatomeeëndieet werd verlaagd door 257 koperstress maar niet door lage atrazineniveaus door de aanwezigheid van een atrazine-258 tolerante, mixotrofe soort (C. closterium). De bijdrage van diatomeeën aan sedimentstabilisatie 259 werd gestimuleerd door context-afhankelijke effecten van beide stressoren. Bij lage 260 atrazineniveaus, verminderden selectieve veranderingen in de gemeenschapsstructuur met 261 dominantie van de atrazine-tolerante maar energie-arme soort *C. closterium*, de dieetkwaliteit 262 met meer dan de helft. Context-afhankelijke stresseffecten alleen beïnvloedden het functioneren van diatomeeën bij hoge atrazine- en koperniveaus. Selectieve en context-afhankelijke 263 264 stresseffecten op dieetkwaliteit beïnvloeden de energietransfer naar het volgende trofische 265 niveau, met M. littorale die de helft van zijn energie-inhoud verliest wanneer gevoed met 266 diatomeeën opgegroeid onder atrazine- en hoge koperstress. Hoofdstuk 5 duidt selectieve 267 stresseffecten aan, die verschuivingen in de gemeenschapsstructuur veroorzaken met een 268 dominantie van soorten met een lage functionele bijdrage, als een meer potente bedreiging voor 269 het functioneren van ecosystemen dan directe stresseffecten op de functionele bijdrage van de 270 soort. De energie-inhoud van de copepoden hing af van die van hun diatomeeënvoedsel, dit 271 onderlijnt de relevantie van de kwaliteit van het dieet als een belangrijke motor van de energie-272 overdracht op de primaire producent en consument interactie.

273 Hoofdstuk 6 integreert de drie hoofdeffecten die ecosysteemfunctionering beïnvloeden onder 274 stress: de functionele impact van selectieve stress, gericht op de functioneel belangrijke soorten, 275 verhoogt met de verschillen in de functionele bijdrage van de soort, en kan worden voorspeld op 276 basis van de respons van de soort en de effectkenmerken. Fysiologische stress, door rechtstreeks 277 verminderen van de functionele bijdrage van de soort, drijft ecosysteemfunctionering door 278 middel van context-afhankelijke effecten bij hoge stressniveaus. Verhogingen van 279 complementariteit, gedreven door de activering van faciliterende mechanismen onder stress, 280 kunnen primaire producenten in staat stellen hun functionele bijdrage te behouden en beperken 281 het verlies van het functioneren van ecosystemen. Door te wijzen op de drie belangrijkste effecten 282 die de diversiteit-functioneringsrelatie onder stress sturen, stelt dit werk een integrale 283 alomvattend kader voor om een betere integratie van antropogene stressoren binnen-diversiteit 284 functionerende onderzoek te garanderen.

#### 286 Chapter 1: General Introduction

#### 287 **1.1. Biodiversity and ecosystem functioning**

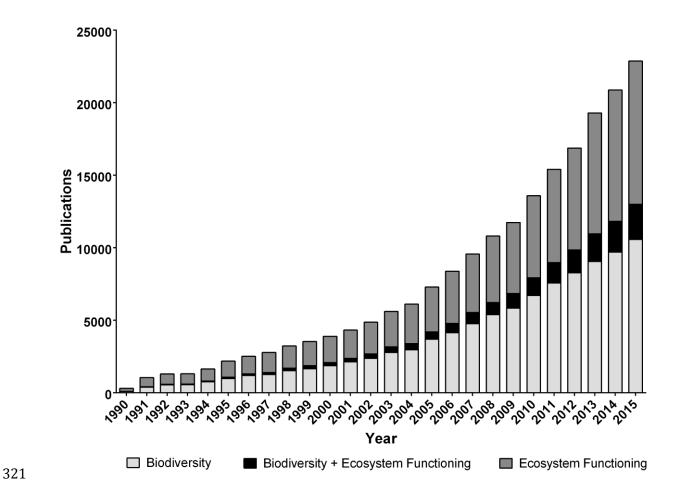
288 Over the past century, Earth has experienced an unprecedented and widespread biodiversity 289 turnover, and current predictions indicate that the rates of biodiversity change will continue to 290 rise in the near future (Sala et al. 2000, Pereira et al. 2010, Barnosky et al. 2011, Dornelas et al. 291 2014, Ceballos et al. 2015, Gonzalez et al. 2016). Human activities have been the main reason for 292 the ongoing extinctions and compositional turnover and will continue to extensively alter 293 biodiversity worldwide (Pimm et al. 1995, Sala et al. 2000, Chapin et al. 2000, Bellard et al. 2012, 294 McGill et al. 2015). Biodiversity has long been treated as a matter of 'pure ecology', which sees 295 biodiversity as a culmination of population and community ecological processes, and changes in 296 biodiversity have sparked mainly aesthetic and ethical concerns (Yarmin 1995, Martín-López et 297 al. 2007, Naeem et al. 2009). This conventional perspective of biodiversity has been 298 revolutionized by a question that had rarely been asked by ecology texts: What is the significance 299 of biodiversity to human wellbeing (Naeem et al. 2009)? This question broadened the 300 conventional perspective of biodiversity as a pure result of ecological processes towards the idea 301 of biodiversity as driver of ecosystem processes, which underpin the ecosystem goods and 302 services that ultimately determine human well-being (Loreau et al. 2001b, Hooper et al. 2005, 303 Srivastava and Vellend 2005, Cardinale et al. 2012, Naeem et al. 2012).

## 304 1.1.1. From community and ecosystem ecology to biodiversity-ecosystem functioning 305 research

306 Until the end of the 20<sup>th</sup> century, the inclusion of biodiversity into ecosystem science has not been 307 a focus in ecological research, which has traditionally been divided into the separate disciplines 308 of community ecology and ecosystem ecology (Loreau 2000, Loreau et al. 2001a). The former 309 focused on how biodiversity and community composition are governed by biotic interactions (e.g. 310 competition, predation, mutualism) and abiotic processes such as disturbance and climate. The 311 latter centred on the properties within ecosystems, such as the rates, dynamics or stability of 312 energy flow or nutrient cycling (Loreau et al. 2001a). In the early 1990s however, the separation 313 of community and ecosystem ecology was overcome when researchers started investigating the 314 impact of biota on ecosystem properties (Lubchenco et al. 1991, Chapin et al. 1992, Hobbs 1992).

Vitousek and Hooper (1993) formally merged community and ecosystem research and proposed
a common positive relationship between biodiversity and ecosystem functioning (hereafter BEF).
BEF research rapidly received widespread attention, and the first BEF experiments investigating
the effects of biodiversity on plant productivity in the mid-1990s (Naeem et al. 1994, Tilman and

- 319 Downing 1994, Tilman et al. 1996) were followed by in a dramatic increase of empirical and
- 320 theoretical studies (Fig. 1.1).



322 Fig. 1.1: The emerging synthesis of biodiversity and ecosystem functioning research. Results represent 323 publications that included either biodiversity (B), ecosystem functioning (EF) or both (BEF, black fill in the 324 centre of bars) in their titles, keywords or abstracts. Note the steep increase in biodiversity research, and 325 the increasing proportion of studies linking biodiversity with ecosystem functioning. Data were obtained 326 from the ISI Web of Knowledge using the 'advanced search' tool and the Science Citation Index Expanded 327 database on 29th July 2016. The general search terms were TS=('biodiversity' OR 'species diversity' OR 328 'species richness' OR 'evenness'\*) for publications on biodiversity, TS=('ecosystem function' OR 'ecosystem 329 functioning' OR 'ecosystem process' OR 'ecosystem') for publications on ecosystem functioning, and TS=('biodiversity' OR 'species diversity' OR 'species richness' OR evenness) AND TS=('ecosystem function' 330 331 OR 'ecosystem functioning' OR 'ecosystem process' OR 'ecosystem') for publications including both terms. 332 The search included the keywords, titles and abstracts of all articles in all languages between 1990 and 333 2015.

- Over the last decade, BEF research has provided an extensive amount of hypotheses and case studies, unraveled the core mechanisms (see 1.6) driving the BEF relation, and allowed for an unambiguous interpretation of the biodiversity effect on ecosystem functioning (Schläpfer and
- 338 Schmid 1999, Loreau and Hector 2001, Loreau et al. 2001b, 2002, Fox 2005, Hooper et al. 2005,
- 339 2012, Cardinale et al. 2011, Tilman et al. 2014). At present, the empirical evidence from more than

340 a thousand BEF studies indicates a positive effect of biodiversity on ecosystem processes such as 341 biomass production, nutrient flux and decomposition (Worm et al. 2006, Balvanera et al. 2006, 342 Cardinale et al. 2006, 2007, 2012, Stachowicz et al. 2007, Finkel et al. 2010, Quijas et al. 2010, 343 Hooper et al. 2012, Tilman et al. 2014, Gamfeldt et al. 2015, Hautier et al. 2015). Although 344 biodiversity effects were by no means universal and the slope of the BEF relation has shown 345 considerable variation, there appears to be a striking level in consistency of biodiversity effects 346 across various organism groups such as microbes, plants and predators, among trophic levels and 347 across the different types ecosystems that have been studied (Balvanera et al. 2006, Cardinale et 348 al. 2006, 2011, Gamfeldt et al. 2008, 2015, Wagg et al. 2014).

#### 349 **1.1.2. Defining biodiversity, ecosystem functioning and ecosystem processes**

350 The term biodiversity encompasses a broad spectrum of biotic scales, which include genetic 351 variation within species, the numbers and relative abundances at the species level, and the 352 diversity of ecosystems on the planet (Purvis and Hector 2000, Mooney 2002, Hooper et al. 2005). 353 This broad scale is reflected in the biodiversity synthesis of the Millennium Ecosystem 354 Assessment (2005), which defines biodiversity as 'the variability among living organisms from all 355 sources including, *inter alia*, terrestrial, marine and other aquatic ecosystems and the ecological 356 complexes of which they are part; this includes diversity within species, between species and of 357 ecosystems.' BEF research commonly focuses on biodiversity at the species level (Loreau et al. 358 2002, Stachowicz et al. 2007), and this thesis consequently describes diversity in terms of both 359 the number of species (species richness) and their relative abundance (species evenness).

360 Although the terms ecosystem process and ecosystem function are often used interchangeably, 361 the term *ecosystem process* refers to the mechanistic processes which are performed by biota (De 362 Groot et al. 2002; in this work e.g. the production of extracellular polymers or fatty acids). These 363 processes regulate *ecosystem functions*, which can be defined as 'the capacity of natural processes and components to provide goods and services that satisfy human needs, directly or indirectly' 364 365 (De Groot et al. 2002; in this work e.g. sediment stabilization or trophic energy transfer). This 366 thesis refers to ecosystem processes when discussing the specific processes carried out by a 367 species or community, and to ecosystem functioning when discussing general concepts of BEF 368 research. Italicised terms are defined in a box after the paragraph where they were first 369 mentioned.

Ecosystem process	Processes performed by biota which regulate ecosystem functioning
Economic functioning	The result of ecosystem processes, providing goods and services that
Ecosystem functioning	satisfy human needs, directly or indirectly

#### 371 **1.1.3. Evenness and Multifunctionality**

372 Most BEF research to date focused on individual ecosystem properties in isolation, with the 373 biomass production of primary producers being the most common functional endpoint 374 (Balvanera et al. 2006, Gamfeldt et al. 2013). Whilst biomass production is vital for delivering a 375 large range of ecosystem services (Imhoff et al. 2004), the contribution of biological communities 376 to decomposition, nutrient cycling, habitat structuring or trophic energy flow are also of key 377 importance for ecosystem functioning (Müller-Navarra et al. 2000, Gamfeldt et al. 2005, Rabaut et 378 al. 2007, Manzoni et al. 2008, Godbold et al. 2009). If diversity effects are not quantified for a larger 379 set of ecosystem functions, BEF research will only partially capture an ecosystem's response to 380 biodiversity loss, since functions other than primary production respond differently to 381 biodiversity change and diversity effects on ecosystem multifunctionality are larger than diversity 382 effects on single functions (Hector and Bagchi 2007, Jiang et al. 2008, Gamfeldt et al. 2013, 383 Lefcheck et al. 2015).

384 Moreover, most BEF studies have chosen species richness as measure of diversity, and have held 385 the relative abundance of species constant across richness treatments (Wilsey and Polley 2004, 386 Hillebrand et al. 2008, Isbell et al. 2009). Until the end of the last decade, 90% of BEF experiments 387 were based on manipulations of species richness, whereas less than 2.5% examined changes in relative species abundances (i.e. species evenness, Balvanera et al. 2006, Hillebrand and 388 389 Matthiessen 2009). However, the impact of a species on ecosystem functioning depends not only 390 on whether it is present in an ecosystem, but also on how abundant it is (Díaz et al. 2003). If the 391 dominant species contributes more to ecosystem functioning than the mean of the community, 392 functioning will increase, and vice versa if the dominant species performs below average 393 (Hillebrand et al. 2008). Human activities do not only alter the number, but also the relative 394 abundance of species (Hillebrand et al. 2008). More importantly, evenness commonly responds 395 more sensitively to environmental change than species richness, and changes in evenness can 396 affect ecosystem functioning long before species are actually threatened by extinction (Chapin et 397 al. 2000, Hillebrand et al. 2008). Thus, the distribution and relative abundance of species within a 398 community might be more important for its contribution to ecosystem functioning than the 399 number of species (Norberg 2004, Hillebrand et al. 2008).

Species richness	The number of species in a community or ecosystem
Species evenness	The relative abundance of species in a community or ecosystem

#### 401 **1.2. Biodiversity and ecosystem functioning in marine systems**

402 Marine ecosystems are of particular interest to conservation science due to their tremendous 403 biodiversity, with a significant number of marine species not yet described, and due to their global 404 significance for the storage and cycling of materials, nutrients, and energy flow (Snelgrove 1999, 405 Covich et al. 2004). Coastal, estuarine and freshwater ecosystems are among the most impaired 406 parts of the biosphere, experiencing some of the highest rates of species loss, but host a 407 disproportionately large fraction of productivity, providing essential services for human well-408 being (Malmqvist and Rundle 2002, Covich et al. 2004, Gamfeldt et al. 2015, Turner and Schaafsma 409 2015). However, most of the early research on the effects of biodiversity on ecosystem functioning 410 has been focussing on terrestrial ecosystems, particularly grasslands, with producer biomass as 411 the predominant functional endpoint (Balvanera et al. 2006, Jiang et al. 2008). Marine and 412 freshwater systems on the other hand have remained a 'niche product' in early BEF research, and 413 conservationists stressed the need to test if conclusions derived from terrestrial experiments also 414 applied to aquatic ecosystems (Covich et al. 2004, Giller et al. 2004). During the last decade, a 415 number of groundbreaking studies responded to these needs and investigated the effects of 416 biodiversity loss on the functioning of marine and particularly coastal systems (e.g. Emmerson et 417 al. 2001, Bolam et al. 2002, Solan et al. 2004, Bracken et al. 2008, Godbold et al. 2009, Vanelslander 418 et al. 2009, Davies et al. 2011, Gustafsson and Boström 2011, Baert et al. 2016). While the majority 419 of BEF research is still based on studies in terrestrial systems (60 % of all BEF contributions as 420 identified by Solan et al. 2009), marine systems are increasingly well represented (11%, the 421 remaining contributions being 13% freshwater studies and 17% generic reviews). By now, the 422 empirical evidence obtained in recent marine studies suggests that biodiversity tends to enhance 423 ecosystem functioning not only in terrestrial, but also in marine habitats (Gamfeldt et al. 2015).

#### 424 **1.3. Anthropogenic stress**

425 In physiological and medical research, the term stress is most generally defined as a condition that 426 causes an organism to shift away from its homeostatic balance (Cannon 1926, Goldstein and Kopin 427 2007, Neumann-Lee 2016). In classic plant ecology, stress has been defined as the reduction in 428 growth or biomass production caused by an external constrain (Grime 1977, Maestre et al. 2009). 429 In an ecosystem functioning context, environmental stress is commonly defined as an 430 environmental condition that impacts ecosystem processes such as biomass production (Mullan 431 Crain and Bertness 2006, Gessner and Hines 2012, Steudel et al. 2012), although this terminology 432 depends on the organism and the range of environments considered (Körner 2003, Mullan Crain 433 and Bertness 2006). A change in environmental conditions which is caused by human activities 434 and affects ecosystem processes can be termed anthropogenic stress (Carlisle and Clements 2005, Halpern et al. 2007, Johnston and Roberts 2009, McMahon et al. 2012, Malaj et al. 2014, NeumannLee 2016).

437 Anthropogenic stress is among the driving forces behind the ongoing biodiversity losses (Lotze et 438 al. 2006, Halpern et al. 2008, Geiger et al. 2010). Among the many anthropogenic impacts on 439 natural ecosystems, chemical pollution is regarded as one of the most potent threats to aquatic 440 biodiversity (Wilcove and Master 2005). In the EU alone there are more than 100 000 registered 441 chemicals (EU 2001), many of which have not been tested for adverse biological effects. Global 442 chemical production is projected to double over the next quarter-century, rapidly outpacing the 443 rate of population growth, and many of these chemicals ultimately enter the environment 444 (Schwarzman and Wilson 2009). The list of chemicals eventually found in coastal ecosystems 445 includes industrial, domestic and agricultural nutrients, pesticides, road run-off, personal care 446 products, disinfectants and pharmaceuticals (European Marine Board 2013), amongst which 447 pesticides and their residues are reported to be among the most devastating agents for aquatic 448 ecosystems, affecting all levels of the food chain from the lowest to the top level (Islam and Tanaka 449 2004). Chemical pollution is jeopardizing biodiversity on a global scale (Wilcove and Master 2005, 450 Backhaus et al. 2012, Malaj et al. 2014), but the percentages of ecological studies addressing 451 pollution do not correspond at all to its ranked importance as a driver for biodiversity loss (Lawler 452 et al. 2006). This underrepresentation of anthropogenic stress in studies testing the effects of 453 biodiversity loss on ecosystem functioning has raised critical questions whether the patterns and 454 diversity-functioning relations obtained in classic BEF experiments reflect those induced by 455 anthropogenic activities (Naeem 2008, Fugère et al. 2012, O'Connor et al. 2015).

Anthropogenic stress A change in environmental conditions which is caused by human activities and affects ecosystem processes

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#### 457 **1.4. Incorporating anthropogenic stress into BEF research**

458 The overwhelming majority of BEF experiments to date have manipulated biodiversity in subsets 459 of species under controlled conditions, which enabled scientists to eliminate functional effects by 460 confounding factors other than biodiversity, and thus allowed an unambiguous interpretation of the BEF relation (Loreau et al. 2001b, Naeem and Wright 2003, Díaz et al. 2003, Srivastava et al. 461 462 2004, Hooper et al. 2005, Cardinale et al. 2006, 2012, Tilman et al. 2014). The high degree of 463 control required in these experiments however implied that the environmental complexity and 464 the anthropogenic drivers that shape BEF relations under global change were largely neglected 465 (Zavaleta and Hulvey 2004, Srivastava and Vellend 2005, Schläpfer et al. 2005, Bracken et al. 2008,

466 Reich et al. 2012, De Laender et al. 2016). The plethora of anthropogenic drivers causing the 467 ongoing biodiversity changes in nature however made it unclear whether the observed BEF 468 patterns will hold for realistic extinction scenarios as induced by anthropogenic change, over 469 multiple trophic levels, and across different types of ecosystems (Giller et al. 2004, Srivastava and 470 Vellend 2005, Raffaelli 2006, Duffy et al. 2007, Hillebrand and Matthiessen 2009, Gamfeldt et al. 471 2013). After the first phase in BEF research had provided the empirical evidence on the direction 472 and the effects underpinning the diversity-functioning relation, the incorporation of realistic 473 biodiversity loss and trophic complexity were thus identified as key challenges for the next 474 generation of BEF research (Hooper et al. 2005, Srivastava and Vellend 2005, Cardinale et al. 475 2012).

#### 476 **1.4.1. The classic BEF design**

477 After early BEF experiments (Naeem et al. 1994, Tilman and Downing 1994) were criticized for 478 confounding species number with species composition, BEF research soon adopted the principle 479 of using random species assemblages to evaluate the functional consequences of changing biodiversity (McGrady-Steed et al. 1997, Hector et al. 1999, Naeem 1999, Tilman et al. 2001, 480 481 Downing and Leibold 2002). To date, most BEF research has involved experiments in which 482 diversity is varied through random draws from a species pool (Srivastava and Vellend 2005, 483 Cardinale et al. 2012, Eisenhauer et al. 2016, Wardle 2016). Random designs have considerably 484 advanced BEF research, by allowing to unequivocally attribute changes in functioning to species 485 richness, and by providing insights into the key mechanisms by which more diverse communities 486 can result in enhanced functioning (Loreau and Hector 2001, Cardinale et al. 2006, Eisenhauer et 487 al. 2016). However, the classic design of random species loss involves three implicit limitations, 488 which are addressed in the following three sections.

#### 489 **1.4.2. Selective stress**

490 First, random species loss implies that the likelihood to be removed from the system is the same 491 for all species, regardless of their extinction-proneness in nature, an assumption which soon drew 492 criticism (Ives and Cardinale 2004, Giller et al. 2004). While it is impractical for experimental 493 studies to mimic all the factors driving species loss in nature, it is clearly an oversimplification to 494 accept random extinction as a representative scenario of species loss (Giller et al. 2004). 495 Anthropogenic stress is likely to cause non-random sequences of biodiversity loss, depending on 496 differences in the species' physiology, morphology, trophic position or habitat specialization, 497 which determine their sensitivity to different stressors (Jonsson et al. 2002, Duffy 2003, Ives and 498 Cardinale 2004, O'Connor and Crowe 2005, Bracken et al. 2008). Stress has therefore been termed 499 selective stress, if it disfavours some species more than others (Wittebolle et al. 2009). This biased diversity loss has raised questions about how useful inferences from randomly assembled
communities will be for conservation efforts, and has led to calls for experimental designs that
remove the most vulnerable species first (Duffy 2003, Raffaelli 2004, Ball et al. 2008, Naeem 2008,
Tilman et al. 2014, Wardle 2016).

#### 504 1.4.3. Physiological stress

505 Second, the same stressors that indirectly alter ecosystem functioning through selective changes 506 in biodiversity can also directly affect the physiology of the surviving species, and thus their 507 contribution to ecosystem functioning. Direct exposure to anthropogenic stressors is however 508 rarely included into the design of BEF studies (McMahon et al. 2012, but see Reusch et al. 2005, 509 Wittebolle et al. 2009, McMahon et al. 2012, Roger et al. 2012, Baert et al. 2016, Radchuk et al. 510 2016). This means that even when diversity gradients are established according to the species' 511 vulnerability in nature, the functional rates of the surviving 'tolerant' species are commonly 512 assumed to remain unaffected. However, stress tolerance may come at a physiological cost, e.g. 513 acclimation by allocating resources from growth to survival pathways (Schimel et al. 2007), and 514 this 'physiological stress' can cause species to contribute differently to a given ecosystem processes 515 than in unstressed conditions (Relyea and Hoverman 2006, Schimel et al. 2007, De Laender et al. 516 2010). This is at least the case for chemical stress, which commonly causes sublethal adverse 517 effects on the surviving species (McWilliam and Baird 2002, Fleeger et al. 2003, Rohr et al. 2006, 518 Liebig et al. 2008). Even when explicitly including exposure to anthropogenic stressors, common 519 BEF designs cannot separate the direct functional impact of a stressor via effects on organismal 520 physiology from its indirect effects that are mediated by changes in biodiversity (Relyea and 521 Hoverman 2006, Radchuk et al. 2016). Disentangling direct stress effects at the physiological level 522 and from indirect effects at the community level thus represents a major challenge to determine 523 functioning in stressed communities (Relyea and Hoverman 2006, Schimel et al. 2007).

#### 524 **1.4.4. Functional contribution**

525 Third, random species loss holds the implicit assumption that all species contribute in a more or 526 less similar way to ecosystem functioning, and that what matters is mainly the number, rather 527 than the characteristics, of species added or lost from a system (Díaz et al. 2003). Most 528 experimental BEF studies have thus documented the 'mean' effect of adding or removing species, 529 regardless of their individual contribution to ecosystem functioning (Lewis 2009). There is 530 mounting evidence, however, that often one or a few particular species have a disproportionate 531 influence on ecosystem properties (Power et al. 1996, Grime 1998, Emmerson et al. 2001, Bolam 532 et al. 2002, Waldbusser et al. 2004, Solan et al. 2004, Bruno et al. 2006, Cardinale et al. 2006). The 533 few studies that investigated selective biodiversity loss caused by anthropogenic disturbance 534 found that anthropogenic stressors can on the one hand have limited effects on functioning 535 despite causing significant diversity loss, as long as the most abundant species are not affected 536 (Smith and Knapp 2003, Solan et al. 2004, Radchuk et al. 2016). On the other hand, BEF studies 537 found a disproportionate loss of functioning when selective stress targeted the species that 538 contributed most to ecosystem functioning (Larsen et al. 2005, McIntyre et al. 2007). Integrating 539 information on the *functional contribution* of a species within scenarios of selective biodiversity 540 loss is thus necessary to successfully predict BEF relationships under stress (Hillebrand and 541 Matthiessen 2009).

542 These three limitations do not devaluate the findings obtained in classic BEF studies which were 543 based on random manipulations of richness under constant experimental conditions. Far from 544 that, in the absence of detailed predictions, these early studies provided a general expectation on 545 how diversity affects ecosystem functioning and unravelled the fundamental mechanisms driving 546 the BEF relations (Hector et al. 2001, Loreau and Hector 2001, Srivastava and Vellend 2005). 547 However, after the incorporation of selective, non-random biodiversity loss was identified as a 548 key challenge for future BEF research, scientists were facing the delicate task to connect the 549 extinction vulnerability of individual species with their contribution to ecosystem functioning 550 (Hooper et al. 2005, Cardinale et al. 2012). The results of random-loss BEF experiments could 551 most easily be used to predict the effects of biodiversity loss when this loss naturally occurs 552 independently of species traits (Srivastava 2002, Lepš 2004, Srivastava and Vellend 2005). 553 However, as discussed above, anthropogenic stress is causing a plethora of non-random scenarios 554 of biodiversity loss, where species are lost according to their sensitivity to a given stressor 555 (McKinney 1997, Srivastava and Vellend 2005, Larsen et al. 2005, Rubach et al. 2012, Malaj et al. 556 2014). This species-specific sensitivity is correlated with the species' biological traits, which can 557 cause organisms to be either sensitive or tolerant to different types of stress (Lavorel and Garnier 558 2002, Solan et al. 2004, Buchwalter et al. 2008, Van den Brink et al. 2011). In their comprehensive 559 review on the future challenges of BEF research, Hooper et al. (2005) thus described 'the 560 covariance between traits affecting extinction and those affecting ecosystem processes' as 'the 561 first critical issue for predicting the consequences of non-random biodiversity loss'.

Functional contribution	The contribution of a species to a given ecosystem process
Dhysiological stross	A change in environmental conditions which affects an ecosystem
Physiological stress	process by altering the species' functional contribution
	A change in environmental conditions which affects an ecosystem
Selective stress	process by selectively targeting species with a high or low
	functional contribution

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#### 563 **1.5. Response and effect traits**

564 The concepts and definitions of 'traits' have varied over time and ecological disciplines (Violle et 565 al. 2007), but BEF research has mainly considered traits as functional traits, which are defined as 566 'morphological, physiological or phenological characteristics of an organism affecting its 567 individual performance' (Violle et al. 2007, Hillebrand and Matthiessen 2009). As such, functional 568 traits are the pillar for the concept of 'functional diversity'. Functional diversity can be described 569 as the diversity of distinct functional groups, in which species are classified according to their 570 traits, or as continuous gradients of different traits, which both allow to predict the changes in 571 ecosystem processes based on changes in community and thus trait composition (Diaz and Cabido 572 2001, Petchey and Gaston 2006, Griffin et al. 2009, Hillebrand and Matthiessen 2009). As such, 573 functional diversity, together with the above-mentioned species richness and evenness, rapidly 574 emerged as third main biodiversity endpoint in BEF research (Balvanera et al. 2006).

575 Suding et al. (2008) and Hillebrand and Matthiessen (2009) introduced a new trait concept into 576 BEF research, which connects response traits (traits that predict species abundance in responses 577 to environmental change) and *effect traits* (traits that reflect the effects of a species on ecosystem 578 processes). This concept corresponded to the need formulated by Hooper et al. (2005), to predict 579 the consequences of selective biodiversity loss by correlating traits predicting extinction with 580 those affecting ecosystem processes. If response traits and effect traits are correlated, i.e. the 581 species contributing most to functioning are also the ones most likely to be lost in response to 582 anthropogenic disturbance, selective biodiversity loss will lead to a disproportionate decline in 583 functioning relative to random biodiversity loss. If response and effect traits are uncorrelated, the 584 community might be able to compensate for the loss of effect traits and maintain its contribution 585 to functioning (Hillebrand and Matthiessen 2009). The presence of toxic chemicals alters not only 586 the abundance of species, but also their functional contribution (McWilliam and Baird 2002, 587 Fleeger et al. 2003, Rohr et al. 2006, but see 1.4.3). For the case of chemical pollution, response 588 traits should thus predict the numerical stress response of species (i.e. changes in abundance, 589 hereafter numerical stress response), and effect traits should furthermore capture the functional 590 stress response (i.e. if species can maintain their functional contribution, hereafter functional 591 stress response). When the numerical and functional stress response are positively related to the 592 same set of response and effect traits, a community decimated by species loss could still be 593 expected to retain most of its functioning. Alternatively, stress exposure could lead to a 594 disproportionate reduction in functioning if numerically tolerant species, while still being able to 595 grow, would lose most of their functional contribution.

596 The use of response traits as predictors of sensitivity to toxic substances has been increasingly 597 advocated in ecotoxicology, as a mechanistic alternative to the otherwise empirical approach of 598 estimating sensitivities using species sensitivity distributions (Baird and Van den Brink 2007, Van 599 den Brink et al. 2011). For example, Baird and Van den Brink (2007) showed that only four 600 response traits could predict the sensitivity of 12 species to a large array of chemical pollutants. 601 Subsequently, several pioneering works have shown the capacity of traits to predict species 602 sensitivity to chemical pollutants, linking traits to specific toxic mechanisms, and thus enabling a 603 more deterministic description of the structure of biological communities under chemical stress 604 (Buchwalter et al. 2008, Guénard et al. 2011, Larras et al. 2012, Rubach et al. 2012, Fischer et al. 605 2013, Pomati and Nizzetto 2013). Whilst the use of response traits is well established, the 606 identification of effect traits and the explicit linkage between both types of traits is still rare in BEF 607 research (but see Pakeman 2011, Heuner et al. 2015, Zwart et al. 2015, Lennon and Lehmkuhl 608 2016), and mostly limited to single species and not in the context of chemical stress. An increased 609 focus on correlating response and effect traits should thus enable scientists to predict how 610 selective changes in diversity caused by environmental change will affect the contribution of 611 biological communities to ecosystem functioning.

Docnonco trait	A trait predicting the abundance of a species or organism group in
Response trait	response to a change in environmental conditions
Effect trait	A trait reflecting the effect of a species or organism group on ecosystem
Ellect trait	processes

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#### 613 **1.6. Complementarity and dominance**

614 The identification of complementarity and selection effects have been among the most crucial findings of early BEF research (Loreau and Hector 2001). After more than two decades of BEF 615 616 research, scientists now widely agree that complementarity and selection are the two key effects 617 which cause diverse communities to perform differently than expected from their monocultures 618 (Hector et al. 2009, Hodapp et al. 2016). The selection effect drives BEF relations through selective 619 processes such as interspecific competition, which can cause dominance (high relative 620 abundance) of species with particular traits. The selection effect can positively or negatively affect 621 a community's contribution to ecosystem functioning, depending on whether the species that 622 ultimately dominate the community have a relatively high or low functional contribution, 623 respectively (Loreau and Hector 2001). In contrast to the competition-driven selection effect, 624 complementarity refers to a class of mechanisms that result in a higher performance of a

625 community than would be expected from the separate performances of each component species 626 (Loreau and Hector 2001). These mechanisms are commonly thought to be facilitation, e.g. if 627 species modify the environment in a way that benefits co-occurring species, and niche 628 partitioning, e.g. different resource requirements among species causing interspecific competition 629 in diverse communities to be lower than intraspecific competition in monocultures (Tilman et al. 630 1997, Loreau and Hector 2001, Duffy et al. 2007, Vanelslander et al. 2009).

631 One limitation of Loreau and Hector's (2001) bipartite partition of the biodiversity effect into 632 selection and complementarity is that it assumes that complementarity is distributed equally 633 across species, i.e. that species mutually benefit each other to the same degree (Hector et al. 2009). 634 However, species with particular traits can perform better than expected in communities either 635 due to competitive replacement of other species, or not at the expense of other species, e.g. by 636 benefitting from complementarity mechanisms without contributing to these mechanisms (the 637 so-called trait-dependent or ,'one-way' complementarity, Fox 2005; Hector et al. 2009). The 638 bipartite partition assigns this one-way complementarity to the selection effect, and may 639 therefore over- or underestimate total complementarity, depending in whether species with 640 particular traits (e.g. high or low monoculture biomass, hereafter yield) benefit more or less from 641 complementarity (Fox 2005; Hector et al. 2009). Therefore, Fox (2005) proposed a tripartite 642 model, that partitions the biodiversity effect on ecosystem functioning into *dominance*, *one-way* 643 complementarity and two-way complementarity effects. The dominance effect occurs when species 644 with particular traits dominate communities at the expense of other species. Depending on 645 whether communities are dominated by high- or low-yield species, the dominance effect will be 646 positive or negative. Trait-dependent or one-way complementarity occurs when species with 647 particular traits perform better in community than in monoculture, but not at the expense of other 648 species. Trait-independent or two-way complementarity occurs when species perform better in 649 community, independently of their traits and not at the expense of other species. Two-way 650 complementarity is thus analogous to complementarity as defined by Loreau and Hector (2001). 651 To date, there have been few explicit comparisons of the bipartite and tripartite partitioning 652 methods (Hector et al. 2009), but where tested, the contribution of one-way complementarity to 653 the total biodiversity effect was minor (Hector et al. 2009, Karlson et al. 2010, Schmidtke et al. 654 2010, Fernandes et al. 2011, Long et al. 2013, Stachova et al. 2013, Siebenkäs et al. 2016). Thus, 655 using the bi- or tripartite partition should produce similar results on the relative importance of 656 dominance / selection and complementarity effects.

More importantly, experimental BEF studies keep producing contrasting results on the relative
contributions of dominance and complementarity effects, leading to an active debate on which
principal effect drives the diversity-functioning relation (Huston et al. 2000, Cardinale et al. 2006,

660 Fargione et al. 2007, Wang et al. 2013, Hodapp et al. 2016). The bottleneck which is limiting the 661 experimental quantification of complementarity and dominance effects, is that in order to 662 unequivocally determine the contribution of either effect to observed BEF relationships, 663 information is required on the yield of every species when grown in community and in 664 monoculture. This information is however largely unavailable (Hector et al. 2009), and as a result 665 complementarity and dominance underpinning observed BEF relationships are largely 666 interpreted indirectly from existing patterns, rather than from direct experimental tests 667 (Cardinale et al. 2006, Hector et al. 2009).

668 The contribution of dominance and complementarity effects to BEF relations under stress is even 669 less clear. Whilst anthropogenic stressors are increasingly incorporated into the design of BEF 670 experiments, these experiments found contrasting biodiversity effects on functioning in stressed 671 communities (clearly positive effects in Mulder et al. 2001, Solan et al. 2004, Zavaleta and Hulvey 672 2004, Larsen et al. 2005, Ives and Carpenter 2007, McMahon et al. 2012, Steudel et al. 2012, Wang 673 et al. 2013, Baert et al. 2016, neutral or even negative diversity effects in (Jonsson et al. 2002, 674 Petchey et al. 2002, Smith and Knapp 2003, Jiang 2007, Fernandes et al. 2011, Roger et al. 2012, 675 Fugère et al. 2012). However, few of these discrepant diversity-functioning relations could be 676 explicitly traced to changes in dominance and complementarity effects (Fernandes et al. 2011, 677 Fugère et al. 2012, Wang et al. 2013, Baert et al. 2016, Hodapp et al. 2016), since quantifying both 678 effects under stress requires data on the yield of every species in community and monoculture at 679 every stress level. Under stressful conditions, species will become dominant by being inherently 680 tolerant and thus replacing other more sensitive species. This is the selective stress described 681 before (see 1.4.2) and although dominance effects could not be explicitly quantified, the functional 682 contribution of these dominant, stress-tolerant species has been identified as key driver of BEF 683 relations under stress (Petchey et al. 2002, Solan et al. 2004, Zavaleta and Hulvey 2004, Larsen et 684 al. 2005, Radchuk et al. 2016). Complementarity effects in stressed communities depend on 685 potential changes in species interactions along stress gradients. Stressful conditions leading to an 686 increase in interspecific competition can cause negative complementarity effects (Fernandes et 687 al. 2011). The complementarity effect increases if facilitation or niche partitioning are enhanced 688 under stress (Fugère et al. 2012; Wang et al. 2013). Direct ecophysiological evidence for 689 interspecific facilitation as driver of complementarity effects is however still limited (Cardinale et 690 al. 2002, Temperton et al. 2007, Vanelslander et al. 2009), and is lacking so far in stressed 691 communities. Determining the relative importance of dominance and complementarity effects as 692 drivers of BEF relations under stress, and connecting both effects to stress tolerance and 693 interactions at the species level, thus remains a largely unresolved task for explaining diversity-694 functioning relations in stressed communities.

Dominance effect	Species with particular traits dominate communities at the expense of
	others.
Two-way complementarity	'Trait-independent' complementarity: growth in community rather than
	monoculture increases the functioning of species, independent of their
	traits and not at the expense of other species.
One-way complementarity	'Trait-dependent' complementarity: growth in community rather than
	monoculture increases the functioning of species with particular traits, but
	not at the expense of other species.

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#### 696 **1.7. Trophic complexity**

#### 697 **1.7.1. Horizontal and vertical biodiversity effects**

698 BEF research has been dominated by experiments focussing on a single trophic level, mostly 699 primary producers in terrestrial grasslands (Petchey et al. 2004, Srivastava et al. 2004, O'Connor 700 and Crowe 2005, Raffaelli 2006, Scherber et al. 2010). Biodiversity loss can act horizontally, i.e. 701 affect the abundance, biomass or resource use at the same trophic level where biodiversity is lost 702 (Cardinale et al. 2006, Duffy et al. 2007, Scherber et al. 2010). However, horizontal species loss 703 may also have vertical consequences, by affecting other trophic levels, organism groups and 704 processes (Cardinale et al. 2006; Duffy et al. 2007; Scherber et al. 2010). Besides the incorporation 705 of realistic non-random species loss, the addition of a vertical dimension of biodiversity was thus 706 soon identified as a major task for BEF research (Hooper et al. 2005, Srivastava and Vellend 2005, 707 Duffy et al. 2007, Reiss et al. 2009, Scherber et al. 2010, McMahon et al. 2012).

708 Although the single trophic level approach has advanced BEF research considerably, it has a 709 number of limitations (Raffaelli 2006, Scherber et al. 2010). For instance, complementarity and 710 dominance effects have been identified as drivers of BEF relations within trophic levels, but these 711 approaches cannot be used to test biodiversity effects across multiple trophic levels, and it is 712 unclear if the same effects occur in multitrophic experiments (Fox 2006, Raffaelli 2006). Another 713 main reason to adopt a second, vertical perspective in BEF research is the need to recognise that 714 biodiversity loss due to anthropogenic activities usually occurs at all trophic levels, and taxa will 715 differ in their sensitivity to particular loss scenarios (Solan et al. 2004, Raffaelli 2006). Thus, 716 anthropogenic stressors can cause differential biodiversity loss and disparate consequences for 717 functioning across trophic levels (Raffaelli 2006, McMahon et al. 2012). There are likely to be 718 significant feedbacks between the trophic level targeted by a stressor and the higher and lower 719 trophic levels (Raffaelli 2006). If these feedbacks are not recognised, the outcomes of onedimensional, single trophic experiments will only be of limited value for policy makers interested

in the impacts of biodiversity loss (Raffaelli et al. 2002, Raffaelli 2006).

722 In principle, adopting a two-dimensional approach links the fields of BEF research (focusing on 723 horizontal diversity and ecosystem functioning) and predator-prey and food web ecology 724 (focusing of vertical diversity), which are thematically linked but have remained largely separate 725 research fields (Duffy et al. 2007). A large number of trophic ecology studies has analysed the 726 mechanisms and importance of trophic interactions (e.g. Pace et al. 1999, Chase et al. 2002, Borer 727 et al. 2005, De Troch et al. 2005, 2012b). Most of this research however has yet to be integrated 728 within BEF research and its (so far horizontal) focus on ecosystem processes (Duffy et al. 2007). 729 After the first generation of BEF research had established the generality of the diversity-730 functioning relation, theoretical and experimental works began to merge the functional effects of 731 diversity and of trophic interactions (e.g. Fox 2004, Petchey et al. 2004, Thebaut and Loreau 2005, 732 Casula et al. 2006). Similar to traditional horizontal BEF approaches, vertical BEF research started 733 in terrestrial environments (Koricheva et al. 2000, Balvanera et al. 2006, Cardinale et al. 2006, 734 Haddad et al. 2009), with vertical BEF efforts in marine systems starting in the second half of the 735 last decade (Gamfeldt et al. 2005, Bruno et al. 2008, Scherber et al. 2010). Recent works found 736 strong correlations between primary producer diversity and ecosystem processes at higher 737 trophic levels, thus confirming the importance of biodiversity in a vertical dimension (Scherber et 738 al. 2010, Lefcheck et al. 2015). These findings also apply to chemical pollutants, which by causing 739 losses of producer diversity, can significantly impact processes at the associated consumer level 740 (Relyea and Hoverman 2006, McMahon et al. 2012). Whilst some of the strongest impacts of 741 biodiversity loss have been found at the plant-animal interface (Scherber et al. 2010, McMahon et 742 al. 2012, Lefcheck et al. 2015), these multitrophic diversity-functioning relationships were less 743 predictable than those found for single trophic levels (Duffy et al. 2007, Reiss et al. 2009, McMahon 744 et al. 2012).

#### 745 **1.7.2. Diet quality**

746 The overwhelming majority of both BEF and trophic ecology experiments to date have focused on 747 *diet quantity* (i.e. the abundance and availability of food, mostly quantified as producer biomass), 748 which indeed is a critical determinant for food web functioning (Arts and Wainmann 1999, 749 Balvanera et al. 2006, Österblom et al. 2008, Scherber et al. 2010, Guo et al. 2016). However, the 750 recent integration of diet biochemistry with traditional studies of diet quantity has indicated that 751 not only abundance and availability of food items, but also *diet quality* can be of striking 752 importance for the reproduction and growth dynamics of many aquatic animals (Arts and 753 Wainmann 1999, Österblom et al. 2008).

754 The differences in the biochemical composition of organisms are most pronounced at the plant-755 animal interface, which features the largest, but also the most variable energy transfer efficiencies 756 in the food web (Brett and Müller-Navarra 1997, Müller-Navarra et al. 2000, De Troch et al. 2012b, 757 Guo et al. 2016). For aquatic primary consumers, algae represent a high quality diet (compared to 758 e.g. terrestrial organic matter), due to their high polyunsaturated fatty-acid (PUFA) content 759 (Taipale et al. 2013, Guo et al. 2016). Fatty acids represent the densest form of energy in aquatic 760 ecosystems (Parrish 2009), and are therefore crucial for the energy flux from primary producers 761 to higher levels of the food web (Brett and Müller-Navarra 1997, Müller-Navarra et al. 2000, Kainz 762 et al. 2004, Gladyshev et al. 2011). PUFAs are defined as fatty acids with more than one double 763 bond, and are almost exclusively generated by plants (Brett and Müller-Navarra 1997) from de 764 novo synthesis of palmitic acid and further enzymatic elongase and desaturation reactions 765 (Harwood and Guschina 2009, Cagliari et al. 2011, Taipale et al. 2013).

766 Amongst all algal classes, Cryptophyceae and Bacillariophyceae (diatoms) are considered as diets 767 of excellent quality for aquatic primary consumers, due to their high content of two specific PUFA 768 components, the so-called essential fatty acids (EFAs) eicosapentaenoic acid (EPA) and 769 docosahexaenoic acid (DHA, Dunstan et al. 1993, Brett et al. 2006, Taipale et al. 2013). Conversely, 770 primary producers with a low concentration of EFAs (e.g. cyanobacteria) are of very poor diet 771 quality for aquatic consumers (Brett et al. 2006, Burns et al. 2011, Taipale et al. 2013). Animals 772 (e.g. crustaceans and fish as well as humans) cannot synthesize these EFAs de novo, and therefore 773 need to obtain these molecules from their diet (Brett and Müller-Navarra 1997, Müller-Navarra 774 et al. 2000, Litzow et al. 2006, Parrish 2009). Diet quality in terms of EFA content is a critical 775 determinant of the growth rate and reproductive success of primary consumers (Müller-Navarra 776 1995, Müller-Navarra et al. 2000, von Elert 2002, Arendt et al. 2005). Moreover, primary 777 consumers such as copepods cannot maintain their energy content when feeding on EFA-poor 778 algae species (Koski et al. 1998, Werbrouck et al. 2016). Algal diet quality can even be a stronger 779 driver of primary consumer biomass than diet quantity (Gladyshev et al. 2011), and a lack of algal 780 EFA production can reduce food web functioning by causing trophic decoupling, i.e. high primary 781 productivity with little corresponding increases in consumer biomass (Guo et al. 2016). Algal EFA 782 production has thus been described as a crucial determinant of diet quality and energy transfer 783 across the food web (Brett and Müller-Navarra 1997, Müller-Navarra et al. 2000, Litzow et al. 784 2006, Parrish 2009, Taipale et al. 2013). The large variation in EFA content among and within 785 algal classes thus makes the diversity and composition of aquatic primary producer communities 786 a key factor for food web functioning (Dunstan et al. 1993, Wyckmans et al. 2007, Taipale et al. 787 2013, Guo et al. 2016).

788 Algal diet quality is strongly influenced by changes in abiotic conditions, and can be altered in 789 response to light intensity (Wainman et al. 1999, Guschina and Harwood 2006), nutrient regimes 790 (Guschina and Harwood 2009, Sanpera-Calbet et al. 2016) and temperature (Morgan-Kiss et al. 791 2006, Piepho et al. 2012, Werbrouck et al. 2016). Far less attention has been devoted to the effect 792 of anthropogenic stress on algal diet quality. Due to the large variation in the EFA content of 793 microalgae (Dunstan et al. 1993, Taipale et al. 2013, Guo et al. 2016), selective changes in algal 794 community composition could potentially change the energy supply for higher trophic levels. 795 Moreover, there is limited but compelling evidence that anthropogenic stressors affect algal EFA 796 production through direct physiological effects. For instance, metal pollution can cause an 797 increasing degree of algal fatty acid saturation (thus lowering the proportion of high-energy EFAs) 798 and a general decrease of EFA production (Chia et al. 2013a, 2013b). Similarly, pesticide stress 799 can cause a general decrease of fatty acid synthesis (Weisshaar et al. 1988, El-Sheekh et al. 1994), 800 or reduce the proportion of EFAs within the total fatty acid pool by inhibiting enzymes involved 801 in fatty acid elongation and thus EFA synthesis (Böger et al. 2000). Nonetheless, diet quality has 802 so far not been integrated in trophic diversity-functioning research. As it has become increasingly 803 clear that diet quality, and not quantity, is the more important factor regulating the efficiency of 804 energy flow through aquatic food webs (Müller-Navarra et al. 2000, Torres-Ruiz et al. 2007, Lau 805 et al. 2009, Gladyshev et al. 2011), the biochemical composition of primary producers should thus 806 be considered when analysing multitrophic BEF relations under anthropogenic stress.

Diet quantity	The abundance and availability of a food source, in this thesis quantified as biomass
Diet quality	The quality of a food source, in this thesis quantified as energy content
Polyunsaturated	Fatty acid molecules with more than one unsaturated carbon bond
fatty acids (PUFAs)	(double bond)
Essential fatty acids	PUFAs which are essential to organismal physiology, which animals
(EFAs)	cannot synthesize and thus have to take up from their diet

807

#### 808 **1.8. Study system and thesis outline**

#### 809 **1.8.1. Study system**

810 This research was conducted in microcosm experiments using benthic diatoms
811 (Heterokontophyta, Bacillariophyceae) as primary producers and harpacticoid copepods
812 (Crustacea, Copepoda, Harpactidoida) as their main consumers, and atrazine and copper as

anthropogenic stressors. Diatoms and copepods were collected at the Paulina intertidal area, a
muddy intertidal habitat in the Westerschelde estuary (SW Netherlands).

815 Within these muddy intertidal systems, microalgal assemblages (microphytobenthos) represent 816 the main primary producers (Underwood and Kromkamp 1999, Forster et al. 2006, Tang and 817 Kristensen 2007). Benthic diatoms are the dominant component of the microphytobenthos, and 818 play an important role for a number of ecosystem processes (Underwood and Kromkamp 1999, 819 Forster et al. 2006, Hicks et al. 2011). Diatoms constitute the main food source for several primary 820 consumers, which makes diatom biomass production essential for the functioning of intertidal 821 food webs (Moens and Vincx 1997, Buffan-Dubau and Carman 2000, De Troch et al. 2012a, Moens 822 et al. 2014, Cnudde et al. 2015). Due to their high essential fatty acid content (EFAs), diatoms are 823 among the most critical components for the energy transfer through marine food webs (Müller-824 Navarra et al. 2000; Taipale et al. 2013; Guo et al. 2016, but see 1.7.2.). Moreover, benthic diatoms 825 excrete large amounts of extracellular polymeric substances (EPS), a mix of high-molecular-weight 826 and carbon-rich molecules (De Brouwer et al. 2000, Smith and Underwood 2000, Underwood and 827 Paterson 2003). EPS play a beneficial role for the diatoms themselves, as a protective measure 828 against valous abiotic disturbances and by enabling diatom locomotion and vertical migration 829 (Pistocchi et al. 1997, Smith and Underwood 2000, Staats et al. 2000, Gerbersdorf et al. 2009b). 830 Moreover, EPS promote ecosystem functioning in marine coastal systems by increasing sediment 831 stability (Decho 1990, 2000, Smith and Underwood 2000, Gerbersdorf et al. 2009a).

832 Meiofauna (organism size between 38 µm and 1 mm) represent an important link between 833 microalgal primary production and higher trophic levels (Montagna et al. 1995, Buffan-Dubau and 834 Carman 2000, Moens et al. 2002, De Troch et al. 2005). Harpacticoid copepods represent an 835 important part of the meiofauna, and are among the main consumers (grazers) of benthic diatoms 836 (Decho and Fleeger 1988, De Troch et al. 2005, 2012a). Their high content of EFAs, which are 837 incorporated directly from their diatom diet, makes harpacticoids an energy-rich food source for 838 small and juvenile fish (Nanton and Castell 1998, Wyckmans et al. 2007). As such, harpacticoid 839 copepods represent a pivotal link for the energy transfer from primary producers to higher trophic 840 levels (e.g. Alheit and Scheibel 1982, De Troch et al. 1998, Buffan-Dubau and Carman 2000, 841 Andersen et al. 2005).

842

Atrazine is a herbicide which binds to the plastoquinone binding protein of photosystem II, causing the disruption of photosynthetic electron flow (Dorigo and Leboulanger 2001, Legrand et al. 2006, Knauert 2008). When atrazine enters freshwater and estuarine environments through flooding events and river runoff, it can have deleterious effects on the growth and photosynthesis of aquatic primary producers, such as microalgae (Peterson et al. 1994, Bester et al. 1995, 848 DeLorenzo et al. 2001, Pennington et al. 2001, Larras et al. 2016). Atrazine is commonly not 849 acutely toxic to aquatic consumers, but has adverse chronic effects on consumers due to food 850 limitation, hormonal disruption and reduced reproduction, although these reproductive and 851 hormonal effects are not consistently observed (Jüttner et al. 1995, Bejarano and Chandler 2003, 852 Bejarano et al. 2005, Forget-Leray et al. 2005, Solomon et al. 2008, Hayes et al. 2011). Despite its 853 Europe-wide ban in 2004, atrazine is still a common pollutant European estuaries and remains 854 one of the most-used pesticide worldwide (Graymore et al. 2001, Carafa et al. 2007, Noppe et al. 855 2007, Benbrook 2016).

856 In contrast to organic pesticides, heavy metals occur naturally in the environment, and several of 857 them are essential for organism physiology (Rengel 1999, Hänsch and Mendel 2009). This is the 858 case for copper which is involved in several metabolic pathways in microalgae, as an essential 859 micronutrient and component of proteins and enzymes (Morelli and Scarano 2004, Hänsch and 860 Mendel 2009). However, copper concentrations above the required levels are toxic to marine 861 organisms at all trophic levels (Welsh et al. 1996, Millward and Grant 2000, Murray-Gulde et al. 862 2002, Real et al. 2003, Manimaran et al. 2012). Copper enters coastal environments through river 863 run-off, industrial and domestic activities, copper mine drainages and agricultural practices, 864 notably its usage in herbicides and antifouling paints (Kennish 1996, Stauber and Davies 2000, 865 Murray-Gulde et al. 2002). Copper affects both primary producers and consumers through the 866 formation of reactive oxygen species (ROS) which can lead to cell death by damaging cell 867 membranes and nucleic acids (Rijstenbil et al. 1994, Knauert and Knauer 2008, Rhee et al. 2013). 868 Copper also affects marine primary consumers by inhibiting membrane transport proteins 869 (Bianchini et al. 2004) and by limiting the quantity of their algal diet (Pinho et al. 2007). Further 870 copper effects on microalgae include the inhibition of electron transport in photosystem II and I 871 (for detailed mechanisms see Miao et al. 2005, Spijkerman et al. 2007), by limiting the reduction 872 of nitrate to nitrite (Manimaran et al. 2012) and reducing the uptake of silicon (in diatoms, Martin-873 Jézéquel et al. 2000).

#### 875 **1.8.2. Thesis outline**

876 The main challenges for future BEF research comprise:

(i) contrasting BEF relations obtained with the classic design of random species loss with thoseinduced by anthropogenic stress (addressed in Chapter 2)

(ii) identifying response and effect traits which predict the functional consequences ofbiodiversity loss under anthropogenic stress (addressed in Chapter 3)

(iii) determining the relative importance of the two main biodiversity effects (dominance and
complementarity) for functioning in stressed communities, and identifying mechanisms which
drive both effects under stress (addressed in Chapter 4)

(iv) analysing the effects of anthropogenic stress on diet quality and food web functioning(addressed in Chapter 5)

886

887 Here, in the general introduction (**Chapter 1**), the scientific setting of this thesis was outlined, 888 with a focus on the recent progress and future research scopes in BEF science. The inclusion of 889 anthropogenic stress into BEF research was identified as first main objective, since it is unclear 890 whether the biodiversity loss and the BEF relations obtained with the classic random BEF design 891 correspond to those induced by anthropogenic stress. Hence, **Chapter 2** contrasts diversity 892 gradients in diatom communities created by the classic design of random community assembly 893 with diversity gradients induced by applying the pesticide atrazine. Next, Chapter 2 tests if BEF 894 relations in both designs are comparable, i.e. if diversity-functioning relationships inferred from 895 random-assembly approaches predict pesticide-driven classic diversity-functioning 896 relationships. Functioning is measured as the diatoms' contribution to four processes: 897 photosynthetic efficiency and biomass production (as proxy for primary production), fatty acid 898 production (as proxy for energy content) and EPS production (as proxy for sediment 899 stabilization). Lastly, Chapter 2 tests if differences between the two types of diversity-functioning 900 relationships could be explained by selective atrazine effects, i.e. to what extent the species' 901 contribution to functioning and their stress tolerance combine in shaping BEF relations under 902 stress.

903 The lacking information on the species' stress tolerance is a main reason why BEF relations are 904 commonly tested with random species loss. **Chapter 3** therefore identifies a set of response and 905 effect traits, which predict the abundance (numerical stress response) and functional contribution 906 (functional stress response) of diatoms under copper and atrazine. By quantifying the effect of atrazine and copper on the abundance (cell densities) and functional contribution (photosynthetic
efficiency, EPS and fatty acid production) of 17 diatom species, Chapter 3 represents a
comprehensive dataset on the effects of two common chemical stressors on benthic diatoms.
Moreover, by linking response and effect traits, Chapter 3 combines ecotoxicology and theoretical
ecology, and provides new insight into the extent to which metal- and herbicide-tolerant species
can maintain their contribution to ecosystem functioning under stress.

913 Dominance (dominance of communities by high- or low-yield species) and complementarity 914 (better performances of species in community due to mechanisms such as resource partitioning 915 or facilitation) are the two main effects which cause communities to perform differently than 916 expected from their component monocultures (see 1.6). However, both effects have rarely been 917 quantified under stress. Moreover, there is little ecophysiological evidence for facilitative 918 mechanisms as driver of complementarity in stressed communities, and BEF models commonly 919 assume complementarity to be mutual, with species reciprocally benefitting each other. **Chapter** 920 **4** tests whether the biodiversity effect on diatom biomass production under copper and atrazine 921 stress depends on dominance or complementarity effects. Moreover, Chapter 4 unravels the 922 fundamental mechanisms driving the diversity effect on functioning, by testing if 923 complementarity under stress is driven by the activation of facilitative mechanisms in stressed 924 diatom communities, and by linking the degree to which species benefit from complementarity to 925 their stress tolerance and trophic mode.

926 **Chapter 5** relates chemical stress effects on primary producers (diatoms) to the consumer level 927 (harpacticoid copepods). First, Chapter 5 measures the effects of atrazine and copper on the 928 contribution of a diatom community to diet quantity (biomass production), diet quality (essential 929 fatty acids production) and sediment stabilization (EPS production). Next, Chapter 5 tests if 930 diatom functioning under stress is driven by changes in community structure or by stress effects 931 on the species' functional contribution. Last, grazer experiments are used to test if atrazine and 932 copper effects on diatom diet quality affect the dominant grazer at the study site, the harpacticoid 933 copepod *Microarthridion littorale*. Thereby, Chapter 5 addresses a main bottleneck issue in 934 current BEF research, which rarely investigates stressor effects across trophic levels and nearly 935 essentially analyzes food web functioning in terms of diet quantity but not diet quality.

936 The main conclusions of this work are summarized and discussed together with future937 perspectives in **Chapter 6**.

938 The **Addenda II to IV** comprise supplementary material to the chapters 2 to 5, respectively.

Addendum V reports the results of an additional experiment which quantified the tolerance of
five harpacticoid species collected at the Paulina intertidal area to direct copper exposure.

30

## 941 Chapter 2: Stressor-induced biodiversity gradients: revisiting 942 biodiversity – ecosystem functioning relationships

943

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948

### 949 Abstract

950 Biodiversity - ecosystem functioning experiments typically inspect functioning in randomly 951 composed communities, representing broad gradients of taxonomic richness. We tested if the 952 resulting evenness gradients and evenness-functioning relationships reflect those found in 953 communities facing evenness loss caused by anthropogenic stressors. To this end, we exposed 954 marine benthic diatom communities to a series of treatments with the herbicide atrazine, and 955 analysed the relationship between the resulting gradients of evenness and ecosystem functioning 956 (primary production, energy content and sediment stabilization). Atrazine exposure resulted in 957 narrower evenness gradients and steeper evenness-functioning relations than produced by the 958 design of random community assembly. The disproportionately large decrease in functioning 959 following atrazine treatment was related to selective atrazine effects on the species that 960 contributed most to the ecosystem functions considered. Our findings demonstrate that the 961 sensitivity to stress and the contribution to ecosystem functioning at the species level should be 962 both considered to understand biodiversity and ecosystem functioning under anthropogenic 963 stress.

964

#### 965 **2.1. Introduction**

966 Biodiversity loss due to human activities has led scientists to enquire how the diversity of 967 communities may regulate ecosystem functioning (Cardinale et al. 2012). During the last two 968 decades, numerous studies have shown that biodiversity favours ecosystem functioning, by 969 positively affecting processes such as productivity or decomposition (Loreau 2000, Hooper et al. 970 2012). Toxic chemicals are potential drivers of biodiversity decrease (De Laender et al. 2014) but 971 remain understudied in biodiversity-ecosystem functioning research and conservation (Lawler et 972 al. 2006). In diversity-functioning research, ecosystem functioning is commonly measured across 973 broad gradients of species richness in randomly assembled communities (Deutschman 2001, Díaz

974 et al. 2003). In the case of toxic chemicals, this makes the implicit assumption that pollution would 975 eliminate a large number of species, and remove individuals in a random way. However, 976 environmentally realistic levels of stressors such as chemical pollution or physical disturbance 977 reduce the abundance of sensitive species (Davies et al. 2011), but without necessarily removing 978 them from the ecosystem, resulting in gradients of evenness rather than species richness 979 (Johnston and Roberts 2009). Whether the evenness gradients created by random manipulations 980 of species richness are representative for those caused by environmental stress such as chemical 981 toxicity remains elusive. Moreover, it is unclear how ecosystem functioning of randomly 982 assembled communities compares to that of communities composed by environmental filtering 983 through chemical toxicity. Chemicals can impact ecosystem functioning directly, by affecting the 984 physiology of the concerned species (De Laender et al. 2010), and indirectly through selective 985 biodiversity loss, depending on the species' sensitivity to the chemical (Vinebrooke et al. 2004). 986 When species differ in their contribution to ecosystem functioning, consequences of biodiversity 987 decline for a particular process can depend on the order in which the species are lost (Solan et al. 988 2004). If the most sensitive species also contributes most to a given ecosystem process, a 989 chemical-induced reduction in density of this species, and therefore evenness reduction, can lead 990 to more pronounced functional consequences than predicted from a random community 991 assembly design.

992 We compare biodiversity gradients and diversity-functioning relations obtained using the design 993 of random community assembly with those generated by filtering through environmental stress. 994 First, we contrast evenness gradients created by randomly assembling 4 benthic diatom species 995 from an intertidal mudflat with those induced by applying the herbicide atrazine, one of the most 996 frequently used herbicides worldwide (Graymore et al. 2001). Despite its EU-wide ban, 997 environmentally critical concentrations of atrazine are still monitored in European intertidal 998 mudflats, which are wetlands of high economic and ecological value (Graymore et al. 2001, Niquil 999 et al. 2006). Next, we test if changes in ecosystem functioning in both sets of experiments are 1000 comparable: Do evenness-functioning relationships inferred from the random assembly approach 1001 predict pesticide-driven non-random evenness-functioning relations? We measured three 1002 ecosystem processes and attributes that are particularly relevant in mudflat ecosystems: primary 1003 production, sediment stabilization and energy content of diatoms. Lastly, we tested if differences 1004 between the two types of diversity-functioning relationships could be explained by selective 1005 atrazine effects, by measuring to what extent species-specific contributions to ecosystem 1006 functioning and species-specific sensitivity to chemicals combine in shaping the diversity-1007 functioning relation.

1008 Diversity-functioning research commonly focuses on the relation between species richness and 1009 primary production (Hillebrand and Matthiessen 2009). Yet, evenness responds faster to 1010 environmental stress and can be a stronger predictor for ecosystem functioning than species 1011 richness per se (Chapin et al. 2000, Kirwan et al. 2007). Also, ecosystem processes other than 1012 primary production can respond differently to diversity loss (Hector and Bagchi 2007). We therefore quantified biodiversity as evenness, and analyzed its relation to three ecosystem 1013 1014 processes and attributes. Biovolume, the most frequently used proxy for phytoplankton biomass 1015 (Hillebrand et al. 1999), and maximum quantum yield of photosynthesis, a proxy for 1016 photosynthetic efficiency (Hartig et al. 1998, Kromkamp et al. 1998), were used to quantify 1017 primary production. Sediment stabilization was quantified as extracellular polymeric substances 1018 (EPS) production. These biopolymers secreted by microphytobenthos and other microorganisms 1019 enhance ecosystem functioning through biogenic sediment stabilization (Gerbersdorf et al. 1020 2009a). The diatom energy content, an attribute that serves as proxy for energy transfer to higher 1021 trophic levels, was estimated by the production of polyunsaturated fatty acids (PUFAs), an 1022 essential diet component for primary consumers (De Troch et al. 2012a). PUFAs are therefore 1023 commonly used as marker for food quality and energy transfer efficiency (Brett and Müller-1024 Navarra 1997, De Troch et al. 2012a).

1025

## 1026 **2.2. Methods**

1027 **Experimental organisms and culture conditions.** Experimental communities were composed 1028 of 4 benthic epipelic diatom species. Navicula arenaria, Nitzschia sp. and Entomoneis paludosa 1029 were sampled from intertidal mudflats in the Westerschelde estuary (SW Netherlands, 51°21'N, 1030 3°43'E). Seminavis robusta was obtained from the culture collection of the Protistology & Aquatic Ecology Research Group (UGent). Cultures of the species were maintained in a climate room at 1031 1032 15±1 °C, a light/dark cycle of 12h / 12h and an illumination of 90 µmol photons m<sup>-2</sup>s<sup>-1</sup>, in culture 1033 medium consisting of filtered and autoclaved natural seawater (salinity  $32\pm1$ ) enriched with f/2 1034 nutrients (Guillard 1975).

1035 **Atrazine and random assembly design.** In the first experiment, diatom communities were 1036 constructed by random community assembly, the common approach in diversity-functioning 1037 research. All possible 15 species combinations were assembled, thereby simulating all 1038 possibilities of random species loss. In the second experiment (atrazine design), diatom 1039 communities were composed selectively through atrazine, by exposing the full species community 1040 to concentrations of 0 (control treatment), 100, 500 and 1000  $\mu$ g/l atrazine respectively. 1041 Technical atrazine (2-chloro-4-ethylamino-6-isopropylamino-*s*-triazine, 99.8% pure) was used in 1042 all treatments. Atrazine was dissolved in the culture medium by means of a 30 min ultrasonic bath. 1043 The resulting atrazine concentrations determined by GC-MS analysis were 91, 465 and 980 µg/l 1044 in the atrazine treatments and  $<1 \mu g/l$  (below detection limit) atrazine in the control treatment. 1045 Treatments in the random assembly design were run twice in polystyrene 6-well-plates, 1046 treatments in the atrazine design in 3 polysterene 6-well-plates: an equivalent of 12 and 18 1047 replicated microcosms per treatment. Each microcosm was inoculated with a total cell density of 1048 10 000 cells/ml (100 000 cells in 10 ml of culture medium), with equal cell densities per species. 1049 Treatments were inoculated with diatom cells from exponentially growing cultures, and 1050 incubated in a climate room at 15±1 °C, under a light / dark cycle of 12h / 12h at 90 µmol photons 1051 m<sup>-2</sup>s<sup>-1</sup>. All experiments were terminated after 18 days.

Biodiversity and ecosystem functioning measurements. Evenness was calculated every second day from cell densities per species. Cell densities were obtained by magnifying and photographing (x100) an area of 0.33 mm<sup>2</sup> per microcosm, using an inverted Axiovert 135 Zeiss microscope (Carl Zeiss, Jena, Germany) and a connected digital camera (Canon PowerShot G11 digital camera). Cell numbers per species were subsequently determined by identifying and counting cells on the computer (ImageJ cell counting software) and converted into cell densities in cells/ml. Evenness for every treatment and replicate was calculated as follows:

1059 
$$J = \frac{H'}{H'_{max}}$$

1060 
$$H' = -\sum_{i=1}^{s} \frac{n_i}{N} \times \log \frac{n_i}{N}$$

Where J is eveness, H' is the Shannon index, n<sub>i</sub> is cell density of the *i*th species. N is total cell density
of the microcosm and H'<sub>max</sub> is log10 of species richness (log10(4) in every treatment). Timeaveraged evenness was used as biodiversity parameter in every treatment.

Biovolume was calculated every second day from cell densities and mean cell biovolume per
species according to (Hillebrand et al. 1999), using measured linear dimensions and formulas
representing the closest approximation of geometric shape for each genus (Addendum I Table S1).
Time-averaged biovolume was calculated as proxy for primary production for every treatment.

1068 Maximum quantum yield of photosynthetic activity was measured at 48-hour-intervals by pulse-1069 amplitude modulated (PAM) fluorometry. Maximum quantum yield is determined as the ratio of 1070 variable and maximum fluorescence ( $F_v/F_m$ ). Maximum fluorescence  $F_m$  is the maximum 1071 fluorescence emission level after a dark adaptation of 20 minutes, measured with a saturating 1072 pulse of light (emission peak at 450 nm, 2700 photons m<sup>-2</sup>s<sup>-1</sup>, 800 ms). Variable fluorescence  $F_v$  is 1073 calculated from the difference between initial fluorescence ( $F_0$ ) and maximum fluorescence ( $F_v = 1074$   $F_m - F_0$ ).

1075 Extracellular polymeric substances were measured by spectrophotometry at the end of the 1076 experiments after 18 days, with 3 replicates of 10 ml suspended diatom culture per treatment. 1077 The suspended diatom cultures were centrifuged for 15 min at 15 °C and 3500g. The supernatant 1078 yielded the soluble EPS fraction, which was left to precipitate overnight at -20°C in 30 ml cold 1079 ethanol (98%), and subsequently centrifuged for 15 min at 15 °C and 3500*g*. The pellet was dried 1080 under a flow of nitrogen and resuspended in 2 ml of 1.5% NaCl. Samples of 200 µl of this 1081 suspension per replicate were used for the EPS analysis. EPS was measured according to a 1082 modified version of the phenol/H<sub>2</sub>SO<sub>4</sub> assay by Dubois et al. (1956). In 24-well-plates, 1 ml 1083 concentrated  $H_2SO_4$  and 200 µl phenol (5%, w / v in distilled water) were added to 200 µl sample. 1084 The mixture was shaken, incubated for 20 min and the absorbance measured at 488 nm using a 1085 Victor<sup>3</sup> multilabel reader (Perkin-Elmer). EPS concentrations were quantified with a function 1086 obtained from a linear regression of known concentrations of glucose standards (ranging from 0.1 1087 to 100  $\mu$ g/ml).

1088 Fatty acid profiles in 2 replicates of 37 ml suspended diatom culture per treatment were analyzed 1089 after 18 days. The suspension was centrifuged for 20 minutes at 4 °C and 2700g. The supernatant 1090 was removed except 5 ml, the pellet was resuspended, transferred into a 10 ml glass tube and 1091 centrifuged for 15 minutes at 4 °C and 2700g. The supernatant was removed and the pellet was 1092 stored at -80 °C until fatty acid extraction. Hydrolysis of total lipid extracts of duplicate diatom 1093 cell suspensions per treatment and methylation to FA methyl esters (FAME) was achieved by a 1094 modified one-step derivatisation method (Abdulkadir and Tsuchiya 2008, De Troch et al. 2012a). 1095 The boron trifluoride-methanol reagent was replaced by 3.5 ml of a 2.5% H<sub>2</sub>SO<sub>4</sub>-methanol 1096 solution per sample since BF3-methanol can cause artefacts or loss of PUFA (Eder 1995). The fatty 1097 acid Methylnonadecanoate C19:0 (Fluka 74208) was added as an internal standard for the 1098 quantification (see further). Samples were centrifuged and vacuum dried. The FAME thus 1099 obtained were analysed using a gaschromatograph (HP 6890N) with a mass spectrometer (HP 1100 5973). The samples were run in splitless mode, with a 1  $\mu$ L injection per run, at an injector 1101 temperature of 250°C using a HP88 column (Agilent J&W; Agilent Co., USA). The oven temperature 1102 was programmed at 50°C for 2 min, followed by a ramp at 25°C min<sup>-1</sup> to 175°C and then a final ramp at 2°C min<sup>-1</sup> to 230°C with a 4 min hold. The FAME were identified by comparison with the 1103 1104 retention times and mass spectra of authentic standards and mass spectral libraries (WILEY, 1105 NITS05), and analysed with the software MSD ChemStation (Agilent Technologies). Quantification 1106 of individual FAME was accomplished by the use of external standards (Supelco # 47885, Sigma-1107 Aldrich Inc., USA). The quantification function of each individual FAME was obtained by linear regression of the chromatographic peak areas and corresponding known concentrations of thestandards (ranging from 5 to 250 ng/ml).

1110 Data analysis. Biodiversity-ecosystem functioning relationships in randomly assembled and 1111 stressed communities were analyzed with a generalized least squares model (Model 1), with the 1112 respective ecosystem process or attribute as response variable and biodiversity as predictor 1113  $(E[EFIa, b] = a + b \cdot B])$ . B represents biodiversity, EF ecosystem functioning, and a and b the 1114 intercept and the slope (i.e. the effect of B on EF). In cases when biodiversity significantly (alpha 1115 = 0.05) predicted ecosystem functioning for both random and stressed treatments, we compared 1116 the slopes (b) and their confidence intervals between the two model fits. The absence or presence 1117 of overlap of the 95% pointwise confidence intervals for b in both models was used as a criterion 1118 for the similarity or difference of diversity-functioning relations between the atrazine-induced 1119 and random biodiversity loss. Normality and homogeneity of model residuals across the evenness 1120 range were inspected using quantile-quantile plots and by plotting residuals versus explanatory 1121 variables, respectively. If homogeneity was violated, the model was refitted using an exponential 1122 variance structure allowing residuals to change with the predictor X (var( $\varepsilon$ ) =  $\sigma 2 \cdot e^{2 \cdot \delta \cdot X}$ ), and 1123 homogeneity was evaluated again (Model 2). By means of likelihood ratio testing, the significance 1124 of these structures was tested (alpha = 0.05).

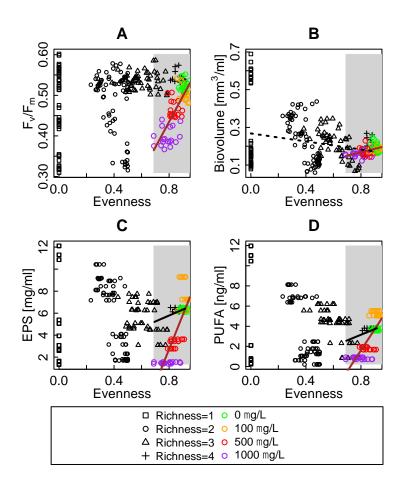
1125 Effects of species identity were assessed by fitting a generalized least squares model to the 1126 respective ecosystem process or attribute as response variable and the cell density of each of the 1127 4 species as predictors [E[EFIa, b] = a +  $\sum_{i=1}^{4} b_i S_i$ ]. EF represents ecosystem functioning, a is the 1128 intercept, S is the cell density of Navicula arenaria, Entomoneis paludosa, Seminavis robusta or 1129 *Nitzschia sp.* respectively, and b represents the slope, i.e. the effect of species identity on EF. 1130 Normality and homogeneity of model residuals was tested as for the diversity-functioning models, 1131 i.e. by plotting residuals vs. all predictors. All calculations were performed in R 2.10.1, using 1132 RStudio (R development Core Team 2015) and the package nlme (Pinheiro et al. 2015).

1133

## 1134 **2.3. Results**

Atrazine reduced diatom evenness, but to a lesser extent than random species removal. Evenness gradients obtained in the atrazine design (0.69 to 0.98) were narrower than those established in the random assembly design (0 to 0.96, Fig. 2.1). Evenness-functioning relationships differed in randomly assembled and atrazine-exposed communities. Evenness was positively related to all 4 ecosystem functioning proxies (all p<0.0001) in the atrazine design (Fig. 2.1, Table 2.1). In the random assembly design, evenness was significantly related to biovolume only (p=0.0006, Fig. 2.1, Table 2.1). Ecosystem functioning was also increasingly variable with decreasing evenness in the random assembly design (Fig. 2.1). This was confirmed by likelihood ratio testing, which showed that diversity-functioning relations in these communities were more accurately described by models including a variance structure allowing the variance of ecosystem functioning to change with evenness (p<0.0001; Table 2.1).

1146



## 1147

1148 Fig. 2.1: Proxies of ecosystem functioning as a function of evenness in randomly assembled and atrazine-1149 exposed diatom communities. Ecosystem functioning is quantified as primary production (photosynthetic 1150 efficiency and biovolume, panels A and B), sediment stabilization (EPS, panel C) and energy content (PUFA, 1151 panel D). Black dots represent diatom communities from the random assembly design, coloured dots denote 1152 diatom communities from the atrazine design. The area marked in grey denotes the evenness gradient 1153 induced by atrazine. Regression lines show significant relations between evenness and functioning. Brown 1154 lines are linear models based on the atrazine design. Dashed and solid black lines are linear models based 1155 on the random assembly design for the whole evenness range or for the evenness range induced by atrazine 1156 (grey area), respectively.

1157

1158	To enable contrasts of evenness-functioning relations induced by the random assembly approach
1159	with those found in stressed communities within the same evenness gradient, we repeated our
1160	analyses, only considering the overlap of both evenness ranges (0.69-0.98) (Fig. 2.1, marked in
1161	grey). This was because the evenness gradient resulting from atrazine exposure was narrower
1162	than that in the random assembly design. These analyses revealed that, in the randomly composed

communities, evenness within a range of 0.69 to 0.98 was positively related to EPS (p=0.02) and PUFA (p=0.02) production (Fig. 2.1, Table 2.1). The slopes of these evenness-functioning relationships however were 3x (EPS) and 6x (PUFA) lower than those found in the atrazine design (Fig. 2.1, Table 2.1), i.e. slopes were less steep at random community composition. Diversityfunctioning relationships were thus different when created by random community assembly than by exposure to atrazine.

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Ecosystem Function	Data	Model	Slope	s.e.	Т	Р	AIC	Log lik.	Validity	LR	P LR
	R	1	0,08	0,02	4,05	0,0001	-416	211	No		
Primary		2	0,09	0,01	4,90	<0,0001	-441	224	No	N.A.	N.A.
Production:	А	1	0,63	0,06	9,84	<0,0001	-264	135	Yes		
(Photosynth.		2	0,63	0,07	8,81	<0,0001	-265	137	Yes	2,62	0,1057
Efficidency)	R'	1	0,03	0,06	0,44	0,6619	-98	52	Yes		
		2	0,04	0,06	0,64	0,5274	-99	54	Yes	N.A.	N.A.
	R	1	-0,11	0,03	-3,01	0,0030	-205	106	No		
Primary		2	-0,11	0,03	-3,50	0,0006	-263	136	Yes	59,82	<.0001
Production:	А	1	0,18	0,04	4,56	<0,0001	-333	170	Yes		
Biovolume		2	0,18	0,04	5,05	<0,0001	-332	170	Yes	0,93	0,3340
[mm <sup>3</sup> /ml]	R'	1	0,33	0,17	1,99	0,0597	-55	30	Yes		
		2	0,42	0,16	2,59	0,0166	-56	32	Yes	3,05	0,0806
	R	1	-0,32	0,81	-0,40	0,6895	904	-449	No		
Sediment		2	0,11	0,74	0,15	0,8835	868	-430	Yes	N.A.	N.A.
stabilization:	А	1	31,12	2,91	10,69	<0,0001	273	-133	Yes		
EPS [mg/ml]		2	29,32	2,63	11,14	<0,0001	274	-133	Yes	1,07	0,3006
[IIIg/IIII]	R'	1	6,43	3,38	1,90	0,0706	81	-38	No		
		2	5,43	2,29	2,37	0,0272	62	-27	Yes	21,60	<.0001
	R	1	-2,36	0,91	-2,60	0,0101	945	-470	No		
Energy		2	-1,05	0,73	-1,44	0,1503	874	-433	Yes	72,77	<.0001
content:	А	1	20,83	2,09	9,94	<0,0001	224	-109	Yes		
PUFA [ng/ml]		2	18,56	1,80	10,30	<0,0001	225	-108	Yes	1,80	0,1794
[118/1111]	R'	1	4,33	3,35	1,30	0,2086	77	-36	No		
		2	6,12	2,38	2,57	0,0175	62	-27	Yes	17,30	<.0001

1171

**Table 2.1:** Relationship between evenness and ecosystem functioning in randomly composed and atrazineexposed diatom communities. Results of generalized least squared models predicting ecosystem functioning from evenness, without (model 1) or with (model 2) a variance structure. 'Data' denotes the data on which the models are based: the classical experiment with random community assembly (R), the atrazine experiment (A), the classical experiment for evenness values overlapping with those from the atrazine experiment (R'); 's.e.' is the standard error on the estimated slopes; T and P denote the t- and pvalues, bold values are statistically significant. AIC is the Akaike information criterion; 'Validity' denotes if residuals were homogeneous and normally distributed ('yes') or not ('no'); 'LR' is the likelihood ratio of

1180 model 1 vs. model 2, P LR the corresponding p-value.

1181

Ecosystem	Data	Duodiatou	Clara		Т	Р	
Function	Data A	Predictor Navicula	Slope -1,31E-01	s.e. 4,66E-02	-2,8243	0,0062	
	A		-1,31E-01 5,08E-03		-2,8243 0,2445	0,8076	
		Entomoneis Seminavis	9,85E-03	2,08E-02 4,29E-02	0,2445 0,2296	0,8078	
Sediment	R'	Nitzschia				<0.0001	
stabilization:			2,57E-01	3,74E-02	6,8742		
EPS [mg/ml]		Navicula	1,94E-02	2,55E-02	0,7579	0,4578	
		Entomoneis	-1,01E-02	8,76E-03	-1,1494	0,2647	
		Seminavis	-4,93E-03	1,31E-03	-3,7631	0,0013	
		Nitzschia	1,83E-02	2,27E-03	8,0803	<.00001	
	А	Navicula	-1,60E+00	1,06E+00	-1,5046	0,1371	
		Entomoneis	1,02E-01	4,69E-01	0,2175	0,8285	
Energy		Seminavis	-1,08E+00	9,70E-01	-1,1102	0,2709	
content:		Nitzschia	6,12E+00	8,50E-01	7,1985	<0.0001	
PUFA [ng/ml]	R'	Navicula	2,93E-01	5,41E-01	0,5429	0,5935	
[8/]		Entomoneis	-2,95E-01	1,77E-01	-1,6690	0,1115	
		Seminavis	-2,12E-01	2,64E-02	-8,0345	<0.0001	
		Nitzschia	4,89E-01	7,29E-02	6,7162	<0.0001	
	А	Navicula	1,73E-01	1,28E-02	13,5532	<0.0001	
		Entomoneis	6,97E-02	6,03E-03	11,5579	<0.0001	
Primary		Seminavis	-8,01E-03	1,24E-02	-0,6463	0,5203	
Production:		Nitzschia	9,59E-02	1,05E-02	9,1230	<0.0001	
Biovolume	R'	Navicula	1,13E-01	1,99E-02	5,6587	<0.0001	
[mm <sup>3</sup> /ml]		Entomoneis	1,46E-01	1,56E-02	9,3537	< 0.0001	
		Seminavis	3,01E-03	2,75E-03	1,0917	0,2886	
		Nitzschia	2,08E-02	1,33E-03	15,6292	< 0.0001	
	А	Navicula	-3,41E-02	3,78E-02	-0,9010	0,3708	
		Entomoneis	2,34E-02	1,96E-02	1,1984	0,2350	
Primary		Seminavis	8,30E-02	4,18E-02	1,9843	0,0513	
Production:		Nitzschia	1,23E-01	3,09E-02	3,9923	<0.0001	
(Photosynth.	R'	Navicula	-2,82E-02	3,85E-02	-0,7319	0,4732	
Efficidency)		Entomoneis	-1,48E-04	1,89E-02	-0,0078	0,9938	
		Seminavis	-4,12E-03	2,87E-03	-1,4325	0,1683	
		Nitzschia	6,53E-03	2,14E-03	3,0548	0,0065	
		-	,	,	,		

1182

**Table 2.2:** Relationship between ecosystem functioning and diatom cell density per species in randomly composed and atrazine-exposed communities. Results of generalized least squared models predicting ecosystem functioning from cell density per species. 'Data' denotes the data on which the models are based: the atrazine experiment (A), or the classical experiment (random community assembly) for evenness values overlapping with those from the atrazine experiment (R'); 's.e.' is the standard error on the estimated slopes; bold values are statistically significant.

1189

1190 As we considered community performance as more relevant for ecosystem functioning than 1191 performance per cell, we initially analyzed diversity-functioning relationships in terms of absolute 1192 values. In order to know if PUFA and EPS production were also related to diversity at the cellular 1193 level, we included supplemental analyses between evenness and PUFA and EPS concentrations 1194 normalized to cell density. Normalized PUFA and evenness were not correlated in the random 1195 assembly design, but positively related in the atrazine design (p<0.0001, Addendum I Fig. S1, 1196 Table S2). EPS production per cell was positively correlated with evenness, both in atrazine-1197 exposed communities and over the whole random diversity gradient (both p<0.0001). Similar to 1198 non-normalized values, the diversity-functioning relation was steeper when evenness gradients 1199 were induced by atrazine. In randomly assembled communities of similar evenness as induced by 1200 atrazine, the relation between evenness and normalized EPS was negative (p=0.02, Addendum I 1201 Fig. S1, Table S2).

1202 In the atrazine design, diversity-functioning relations were steeper, although diversity loss was 1203 less severe than predicted by the random assembly approach. In the random assembly design, 1204 *Nitzschia sp.*, if present, always became the dominant species. Exposure to atrazine did not lead to 1205 species removal, but to a change in dominance: instead of Nitzschia sp., Navicula arenaria 1206 dominated communities exposed to 500 and 1000  $\mu$ g/l atrazine, reaching its highest cell density 1207 at the two highest atrazine concentrations. Detailed cell densities per species for every treatment 1208 are indicated in Addendum I Fig. S2 for the random assembly design and Addendum I Fig. S3 for 1209 the atrazine design. Moreover, *Nitzschia sp.* cell density was positively correlated with all proxies 1210 of ecosystem functioning, in both atrazine-exposed and randomly composed communities (Table 1211 2.2). PUFA and EPS production, the processes that were related to evenness in both the atrazine 1212 and the random assembly design, thus also correlated with the abundance of Nitzschia sp. 1213 Evenness gradients found in atrazine and random assembly treatments hence coincided with a 1214 change in the identity of the dominant species (dominance of Navicula arenaria and Nitzschia sp. 1215 respectively). At high atrazine concentrations the growth of *Nitzschia sp.*, the species contributing 1216 most to all analyzed functions, was suppressed. Instead, communities were dominated by 1217 Navicula, which was negatively correlated with EPS production in the atrazine-exposed 1218 communities (Table 2.2).

1219 The observed diversity-functioning relationships were thus related to the identity of the most 1220 productive species, and a selective suppression of this species by atrazine can explain the steeper 1221 decline of functioning at stressor-induced than at random biodiversity loss.

1222

#### 1223 **2.4. Discussion**

1224 In biodiversity-ecosystem functioning research, one typically aims at experimental biodiversity 1225 gradients that are as broad as possible, to test the effects of widespread diversity loss on 1226 ecosystem functioning (Naeem 2008). Indeed, narrow, evenness-based biodiversity gradients are 1227 uncommon in experimental diversity-functioning research (Hillebrand et al. 2008). Instead, 1228 biodiversity loss by anthropogenic activity is typically mimicked with broad diversity gradients, 1229 covering the entire range from minimal to maximal species richness of the considered ecosystem 1230 (Hillebrand et al. 2008), corresponding to evenness ranges from 0 to 1. In the case of toxic 1231 chemicals however, such extensive biodiversity loss is unlikely to happen. Realistic biodiversity 1232 loss most often only affects a fraction of the existing species pool, resulting in alterations of 1233 evenness rather than richness (Pennington et al. 2001, Hillebrand and Matthiessen 2009), even at 1234 chemical concentrations well above environmental concentrations, as represented by the highest 1235 atrazine concentrations in our study (Pennington et al. 2001). Evenness alterations induced by 1236 atrazine were less severe than those in the richness-based random assembly approach, which in 1237 this case overestimated the biodiversity loss induced by chemical stress. Yet in the random 1238 assembly design, we recorded positive effects of diversity on energy content and sediment 1239 stabilization within a narrow range of high evenness, but not over the whole broad evenness 1240 gradient. Rather than being a tradeoff to realism, we thus suggest narrow biodiversity gradients, 1241 with evenness instead of richness as diversity proxy, as a more realistic simulation of pollution-1242 induced biodiversity loss that can facilitate the detection of biodiversity effects on ecosystem 1243 functioning.

1244 Diversity-functioning relations were steeper when diversity gradients were chemically-induced 1245 rather than randomly assembled, denoting a disproportionately large decrease of functioning in 1246 diatom communities under stress conditions. Contrasting the design of random community 1247 assembly with scenarios of selective biodiversity loss due to environmental filtering is among the 1248 main challenges for future diversity-functioning research (Srivastava 2002, Giller et al. 2004), but 1249 has rarely been tested experimentally (Symstad and Tilman 2001, Smith and Knapp 2003, Solan 1250 et al. 2004, Larsen et al. 2005). Chemical stress moreover has not been studied as driver for 1251 realistic biodiversity loss and ecosystem functioning, although toxic chemicals emitted by 1252 industry, agriculture and households are among the most potent threats to aquatic species (Geiger 1253 et al. 2010). We found a larger discrepancy in random and stressor-induced diversity-functioning 1254 relations (3 to 6 time steeper relations in stressed communities in terms of EPS and PUFA 1255 production) than in other cases where the random assembly design could be contrasted with 1256 realistic diversity loss (Smith and Knapp 2003, Solan et al. 2004, Schläpfer et al. 2005, Larsen et 1257 al. 2005). This can be due to the organisms and ecosystem processes studied, and to selective 1258 atrazine effects on the physiological rates and composition of the diatom communities.

1259 In our diatom communities that were randomly assembled, primary production, the most 1260 common measure of ecosystem functioning (Hooper et al. 2005), was not related to diversity. In 1261 contrast, sediment stabilization and energy content showed a steeper relation to diversity than 1262 primary production in both the atrazine and the random assembly design, which facilitated the contrast of the two experimental approaches. The observed functional consequences of stressor-1263 1264 induced diversity loss thus depended on the measured ecosystem function, which underlines the 1265 value of analyzing multiple ecosystem processes (Hector and Bagchi 2007, Hiddink et al. 2009). 1266 Also, the pool of 4 species could have contributed to the steep diversity-functioning relations. In 1267 our experimental communities, the functionally most important species was most affected by 1268 atrazine (but see below). In a larger species pool, this selection effect could be compensated by 1269 functionally redundant, but less sensitive species. Moreover, as randomized and stress-induced 1270 biodiversity and ecosystem functioning have so far not been assessed in marine microalgae, we 1271 cannot relate our findings to other studies using similar organisms. The disproportionate loss of 1272 functioning following selective loss of diatom biodiversity however corresponds to existing 1273 knowledge on non-random diversity loss in grassland primary producers (Symstad and Tilman 1274 2001, Zaveleta and Hulvey 2004). Yet, the steep diversity-functioning relationships in our stressed 1275 communities can not only be explained by the number and type of different organisms and 1276 ecosystem processes. Loss of functioning in stressed communities can be related to selective 1277 alterations of biodiversity by the stressor, as well as direct stressor effects (Schimel et al. 2007). 1278 In our case, atrazine could affect functioning indirectly through selective changes of community 1279 composition and directly by reducing physiological rates of the species. Atrazine treatments in 1280 which functioning was reduced with regard to the control were essentially dominated by Navicula 1281 arenaria, the only species which was able to grow at the highest atrazine concentration. To inspect 1282 potential direct physiological effects, we compared PUFA and EPS production per cell in these N. 1283 arenaria-dominated stressed communities with those of unexposed N. arenaria monocultures. 1284 Cells in the *N. arenaria*-dominated communities produced 1.8 and 3.5 times less EPS, but 2 and 3 1285 times more PUFA than in the monocultures of this species (Addendum I Fig. S4). The reduction in 1286 EPS in stressed communities could thus be due to physiological effects of atrazine on *N. arenaria*. 1287 The reduction in PUFA however does not seem to be caused by physiological effects, as *N. arenaria* 1288 actually seems to increase its PUFA production when exposed to atrazine. To understand the steep 1289 loss of functioning in our stress-exposed communities, we must therefore, depending on the 1290 ecosystem process, look beyond direct stressor effects and consider selective atrazine effects on 1291 the community composition.

In diversity-functioning research, consequences of biodiversity loss are commonly predicted
using randomly assembled communities, whereas environmental filtering due to stress results in
distinct non-random community compositions. Furthermore, evenness is often held constantly

1295 high across species richness treatments (Wilsey and Polley 2004), thus minimizing effects of 1296 relative abundances (i.e. species identity) on ecosystem functioning. Due to the increasing 1297 recognition that biodiversity and species identity combine in shaping ecosystem functioning 1298 (Bruno et al. 2006), research on realistic biodiversity loss has started by removing rare and 1299 uncommon species from the community (Smith and Knapp 2003). Despite making the step from 1300 random to a more realistic, abundance-based biodiversity loss, this approach still assumes that 1301 the naturally abundant species are also most resistant to environmental stressors and less likely 1302 to be lost. In our diatom communities, Nitzschia sp. was the most abundant species in the natural 1303 samples as well as in the experimental communities, contributing most to the measured 1304 ecosystem processes. However, when exposed to stress, Nitzschia sp. was the species whose cell 1305 density was most reduced by atrazine. This shows that both in stressed and randomly assembled 1306 communities, community composition tends to be uneven, and that the most abundant species is 1307 not necessarily the most stress-resistant. Biodiversity loss generally reduces ecosystem 1308 functioning, but general physiological and selective stressor effects on the most abundant and 1309 functionally most efficient species lead to a disproportionate loss of ecosystem functioning in 1310 stressed communities compared the random assembly design.

1311 Benthic diatoms are the dominant primary producers and food source in many intertidal, soft-1312 sediment environments as well as on rocky shores (Tang and Kristensen 2007, Speybroeck et al. 1313 2008). Diatom communities play a pivotal role in controlling energy flow, nutrient fluxes and 1314 productivity in these systems (Underwood and Kromkamp 1999, De Troch et al. 2012a). Diatom 1315 biodiversity loss resulting in a two-fold reduction in diatom biomass, three-fold reduction of 1316 unsaturated fatty acids and six-fold decrease of EPS production would strongly impact the 1317 ecosystem under concern. The standing stock and energy flux to higher trophic levels would be 1318 affected (Decho 1990, Müller-Navarra et al. 2000), zoobenthos and as a consequence fish biomass 1319 production limited (De Troch et al. 2012a), and sediment erosion promoted. Concomitantly, 1320 knock-on effects on the variety of processes undertaken by sediment-associated organisms 1321 collectively contributing to ecosystem functioning can be expected (Tolhurst et al. 2003). 1322 Stressor-induced biodiversity loss at the base of the marine food web could thus affect ecosystem 1323 functioning more seriously than predicted by the random assembly design in diversity-1324 functioning research. In our study, the random assembly approach overestimated biodiversity 1325 loss, but underestimated the associated loss of ecosystem functioning. These contrasting results 1326 strengthen the need to more realistically predict biodiversity gradients and diversity-functioning 1327 relations caused by anthropogenic stress, by considering the prevailing stressors, and the 1328 sensitivity and contribution to ecosystem functioning of the concerned species.

1329 In conclusion, we infer three general statements from our results. First, existing diversity-1330 functioning research largely overestimates the biodiversity gradients induced by common 1331 environmental stressors such as herbicides. Second, diversity-functioning relations differ 1332 depending on whether the diversity gradient is generated by random community assembly, or by 1333 stressor-induced non-random assembly. Third, physiological stressor effects can affect ecosystem 1334 functioning, and the slope of the diversity-functioning relation depends on stressor-induced 1335 dominance shifts.

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1344

## 1345 Chapter 3: Different response-effect trait relationships underlie 1346 contrasting numerical and functional responses to two chemical

## 1347 stressors

1348

1349 Under revision:

Mensens, C., De Laender, F., Janssen, C. R., Sabbe, K & De Troch, M. (2016) Different response-effect
trait relationships underlie contrasting numerical and functional responses to two chemical
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1353

## 1354 Abstract

Environmental stress can impact ecosystem functioning through the loss of species, but also by altering their densities and functional contribution. Our understanding of the capacity of species to withstand stress effects on density ('numerical response') and on functional contribution ('functional response'), and how these are related to one another, is limited, which hampers efforts to predict ecosystem functioning under stress.

We first identified the response traits and effect traits, driving the numerical and functional response respectively (potential contribution to primary production, sediment stabilization and energy content) of 18 marine benthic diatom strains to two different chemical stressors (a pesticide and a metal). We then examined if the functional contribution of diatoms in unstressed conditions is related to their numerical and functional stress response. Finally, we investigated the relationship between numerical and functional response, i.e. whether numerically tolerant species could maintain their contribution to ecosystem functioning under stress.

The response-effect trait relationship was different depending on stressor type: diatom numerical and functional response (for all three functions) to the metal were predicted by the same set of intercorrelated morphological traits, with large cells being more stress-resistant. Numerical and functional response to the metal were positively related, with species whose density was least affected also maintaining most of their functional contribution. These species also contributed most to primary production and sediment stabilization in unstressed conditions.

1373 In contrast, the capacity for mixotrophic growth predicted the numerical, but not the functional 1374 response to the pesticide. The relation between numerical and functional response to the 1375 pesticide was positive for primary production, but negative for energy content and sediment 1376 stabilization, with numerically tolerant species being most affected in their functional 1377 contribution. Synthesis. Ecosystem functioning loss following exposure to stress can be predicted from response and effect traits, which drive the species' numerical and functional stress response. The extent of functional loss will depend on the relation of both types of traits, with a disproportionate loss of functioning if traits predicting numerical response are not related to those driving functional response.

1383

### 1384 **3.1. Introduction**

1385 Understanding and predicting the consequences of anthropogenic stress on the functioning of 1386 ecosystems is becoming increasingly critical in view of the profound pressure of human activities 1387 on natural ecosystems and the goods and services they deliver (Balvanera et al. 2006, Halpern et 1388 al. 2008). Environmental stress can affect ecosystem functioning by causing species loss, but also 1389 by altering their densities and contributions to functioning (Fox 2006). While the effects of species 1390 loss on ecosystem functioning have been extensively documented (Naeem et al. 2012), realistic 1391 scenarios which also incorporate changes in species densities and functional contributions are 1392 still rare, which hampers realistically predicting the overall effect of environmental stress on 1393 ecosystem functioning (Duffy et al. 2007, Hooper et al. 2012).

1394 Community structure along environmental gradients has been related to the morphological, 1395 biochemical and physiological traits of the species present (Litchman et al. 2007, Violle et al. 1396 2007). Consequently, trait-based approaches are promising to predict species sensitivities to 1397 different types of stress (Baird and Van den Brink 2007, Larras et al. 2012). Suding et al. (2008), 1398 Hillebrand and Matthiessen (2009) and Litchman et al. (2015) proposed a trait-based framework 1399 to evaluate ecosystem functioning under global change which incorporates the relationship 1400 between traits predicting species densities under stress (response traits, which affect the 1401 numerical stress response of species) to traits affecting ecosystem functioning (effect traits, which 1402 affect the functional stress response of species). This is essential as correlations between response 1403 and effect traits can strongly modify the impact of stress on ecosystem functioning. If the 1404 numerical and functional stress response are positively related to the same set of response and 1405 effect traits, a community decimated by species loss could still be expected to retain most of its 1406 functioning. Alternatively, stress exposure could lead to a disproportionate reduction in 1407 functioning if stress-resistant species, while still being able to grow, would lose most of their 1408 functional contribution. Stressor effects on ecosystem functioning would then be more important 1409 than merely estimated from species densities. Experimental approaches addressing this 1410 relationship are still rare (Hillebrand and Matthiessen 2009, but see Heuner et al. 2015), and 1411 identifying traits which generally predict stressor effects on a taxon's numerical and functional 1412 stress response would be a key step towards predicting community composition and functioning 1413 under different types of stress. Toxic chemicals are among the most common abiotic stressors. 1414 They cause widespread loss of biodiversity and ecosystem functioning (Malaj et al. 2014) but 1415 remain understudied in conservation science (Lawler et al. 2006). The numerical and functional 1416 response to chemical stress has only been quantified for a few species and toxicants, because 1417 generating these data is time- and labour-intensive (Larras et al. 2014). As a result, in 1418 ecotoxicological and also ecological research in general, species loss is often assumed to be 1419 random (O'Connor and Crowe 2005, Baird and Van den Brink 2007, Mensens et al. 2015), and 1420 contributions of species to ecosystem functioning are considered to be equivalent (Díaz et al. 1421 2003). However, species tend to differ in their numerical and functional stress response, as toxic 1422 substances do not only alter species densities, but may also impose a physiological cost, as such 1423 affecting their contribution to ecosystem functioning (Schimel et al. 2007, De Laender et al. 2010).

1424 In the present paper, we relate the traits driving the numerical and functional response of 18 1425 benthic diatom strains to metal (copper) and pesticide (atrazine) stress. Benthic intertidal 1426 diatoms play a pivotal role in the functioning of estuarine ecosystems, where they can provide up 1427 to 50% of primary production (Underwood and Kromkamp 1999), enhance sediment stabilization 1428 through the secretion of extracellular polymers (Decho 2000), and are of essential dietary 1429 importance for higher trophic levels (Dunstan et al. 1993). Copper is one of the most common 1430 pollutants in coastal ecosystems, due to input through urban and industrial runoff and its use in 1431 herbicides and antifouling paints (Kennish 1996; Murray-Gulde et al. 2002). Atrazine is one of the 1432 most frequently used herbicides worldwide (Graymore et al. 2001) and represents a common 1433 stressor in intertidal mudflats, which are wetlands of high economic and ecological value 1434 (Graymore et al. 2001; Niquil et al. 2006). Atrazine and copper are both reported to affect diatom 1435 cell densities and functioning by different modes of action: atrazine blocks the electron transport 1436 chain of Photosystem II (PSII), resulting in the excitation energy generated in PSII being lost as 1437 heat or fluorescent light instead of being used for CO<sub>2</sub> assimilation in the Calvin cycle (Dorigo and 1438 Leboulanger 2001). An excess of copper decreases the uptake of silicon, inhibits the reduction of 1439 nitrate to nitrite and leads to the production of reactive oxygen species that damage cell 1440 membranes, nucleic acids and chloroplasts (Knauert and Knauer 2008; Masmoudi et al. 2013).

We first test if the numerical and functional response to copper and atrazine in diatoms can bepredicted from a set of morphological, physiological and biochemical traits.

Next, we analyze the relation between the functional contribution in unstressed conditions and
the numerical and functional response to atrazine and copper, for instance if the species with the
highest functioning (e.g. contributing most to primary production or energy flow in intertidal
systems) tend to be particularly tolerant or sensitive to either of the stressors.

Last, we test if and how the numerical and functional response are related, for example if species
that are tolerant in terms of density (i.e. their growth is unaffected) are also able to maintain their
functional contribution under chemical stress.

1450 The functional contribution of each diatom strain was quantified as the contribution per cell to 1451 three processes that are crucial for ecosystem functioning in intertidal systems. The potential 1452 contribution to sediment stabilization was estimated as extracellular polymeric substances (EPS) 1453 production. These biopolymers secreted by microphytobenthos enhance ecosystem functioning 1454 through biogenic sediment stabilization (Gerbersdorf et al. 2009a). Maximum light use efficiency 1455 in photosystem II (hereafter photosynthetic efficiency), an essential component in primary 1456 production, was measured as maximum quantum yield of photosynthesis in photosystem II 1457 (Cleveland et al. 1989, Consalvey et al. 2005, Barnett et al. 2015). The diatom energy content for 1458 higher trophic levels was quantified as the production of polyunsaturated fatty acids, an essential diet component for primary consumers and a marker for energy transfer efficiency (Brett and 1459 Müller-Navarra 1997; De Troch et al. 2012a). 1460

To predict numerical and functional stress response, we selected morphological and biochemical traits (biovolume, surface-to-volume ratio, cell length, total organic carbon, total nitrogen and chlorophyll a content), which have been linked to the functional properties and response to environmental change in microalgae (Finkel et al. 2010, Kruk et al. 2010). Furthermore we tested if the capacity for mixotrophic growth, which explained diatom sensitivity to herbicides in recent studies (Debenest et al. 2009, Larras et al. 2014), could explain the numerical and functional response of diatoms to chemical stress.

1468

#### 1469 **3.2. Methods**

1470 Experimental organisms and culture conditions. In total 18 diatom strains (belonging to 17 1471 species, Addendum II Table S1) were isolated from intertidal mudflats in the Westerschelde 1472 estuary (The Netherlands, 51°21'N, 3°43'E) and the Veerse meer (The Netherlands, 51°32'N, 1473 3°47'E), where diatoms represent the dominant component of the microphytobenthos (Forster et 1474 al. 2006). The strains were deposited in the diatom culture collection (BCCM/DCG) of the Belgian 1475 Coordinated Collection of Micro-organisms (http://bccm.belspo.be); accession numbers are 1476 indicated in Addendum II Table S1. Diatoms were maintained in a climate room at 15±1 °C, a 1477 light/dark cycle of 12h / 12h and an illumination of 90 µmol photons m<sup>-2</sup>s<sup>-1</sup>, in culture medium 1478 consisting of filtered and autoclaved natural seawater (salinity  $32\pm1$ ) enriched with f/2 nutrients 1479 (Guillard 1975).

1480 Atrazine and copper experiments. Diatoms were exposed to concentrations of 0, 50, 100, 200, 1481 500 and 1000  $\mu$ g/l atrazine and 0, 10, 30, 90, 270 and 810  $\mu$ g/l copper respectively. 1482 Concentrations were based on trial tests (data not shown) as well as published sensitivity data for 1483 marine benthic diatoms (Pistocchi et al. 1997, Levy et al. 2007, Wood et al. 2014) and were 1484 prepared from atrazine and copper stock solutions. The atrazine stock solution was prepared by 1485 dissolving 50 mg technical atrazine (2-chloro-4-ethylamino-6-isopropylamino-s-triazine, 99.8% 1486 pure, Sigma-Aldrich Chemie Gmbh, Munich, Germany) in 10 ml acetone as a carrier to increase the 1487 solubility of atrazine, with a maximum final volume of 0.025% acetone in the treatments. An 1488 acetone control treatment of 0.025% acetone was included and compared to an acetone-free 1489 control to test for carrier effects. All atrazine treatments were compared to the acetone control. 1490 Copper (as a Cu[II]Cl<sub>2</sub> solution, analytical grade; VWR International) was spiked directly into the 1491 culture medium before exposure of the diatoms. F/2 culture medium for the copper experiments 1492 was prepared without EDTA, to avoid complexation of free copper ions (Pistocchi et al. 1997). The 1493 resulting atrazine and copper concentrations are indicated in Addendum II Table S2. All 1494 experiments were run in polysterene 6-well-plates, an equivalent of 6 replicated microcosms per 1495 treatment. Each microcosm was inoculated with a total cell density of 5000 diatom cells/ml (50 1496 000 cells in 10 ml of culture medium) from exponentially growing cultures, and incubated in a 1497 climate room at 15±1 °C, under a light / dark cycle of 12h / 12h at 90 µmol photons m-2s-1. Culture 1498 medium spiked with atrazine and copper respectively was refreshed after 12 days, and all 1499 experiments were terminated after 25 days.

1500 Cell density, traits and functional contribution. Cell densities were obtained on days 0, 2, 5, 10, 1501 15, 20 and 25 by magnifying and photographing (x100) an area of 0.33 mm<sup>2</sup> per microcosm, using 1502 an inverted Axiovert 135 Zeiss microscope (Carl Zeiss, Jena, Germany) and a connected digital 1503 camera (Canon PowerShot G11 digital camera). Cell numbers (cell densities in cells/ml) were 1504 subsequently determined by digitally counting the cells by means of ImageJ cell counting software 1505 (Schneider et al. 2012).

Morphological traits (biovolume, length, surface to volume ratio) were obtained by
photographing diatom cells in every replicate, using an inverted Axiovert 135 Zeiss microscope
(Carl Zeiss, Jena, Germany) and a connected digital camera (Canon PowerShot G11 digital camera).
The linear dimensions were digitally measured using ImageJ software calibrated to a stage
micrometer and a standardized set of equations representing the closest approximation of
geometric shape for each genus (Hillebrand et al. 1999).

1512 Chlorophyll a content was measured *in vivo* on days 0, 2, 5, 10, 15, 20 and 25 of the experiments
1513 by PAM fluorometry (MAXI Imaging PAM fluorometer, Walz, Germany) as the minimum
1514 chlorophyll fluorescence signal (F<sub>0</sub>) after 15 minutes of dark-adaptation.

Total organic carbon (TOC) and total nitrogen (TN) were measured in a separate experiment,
under the same conditions as the atrazine and copper experiments described above. After 25 days
two replicates of 10 ml suspended diatom culture per treatment were harvested and centrifuged
10 minutes at 4 °C and 1000*g*. The pellets were freeze-dried, homogenized and acidified with 1%

- 1519 HCl and TOC and TN content were measured using a Flash EA 1112+ MAs 200 elemental analyser
- 1520 (Thermo Interscience).
- 1521 Mixotrophy was measured in a separate experiment as the capacity to compensate growth 1522 inhibition by atrazine in the presence of organic carbon sources. Each diatom strain was grown in 1523 4 treatments: (a) f/2 culture medium, (b) f/2 medium + atrazine (at EC50 concentration), (c) f/21524 medium + atrazine + 0.1 mM Glucose (d) f/2 medium + atrazine + 0.6 mM Glucose. Mixotrophy 1525 was assessed as the ratio of the maximum cell density (carrying capacity K) that each strain 1526 reached under atrazine (treatment b) and atrazine + glucose (treatments c and d): K(c)/K(b) and 1527 K(d)/K(b). The methodology and glucose concentrations of this experiment were set as described 1528 in Lewin and Hellebust (1975). Each microcosm was inoculated with a total cell density of 5000 1529 diatom cells/ml, cell densities were measured on days 0, 5, 10, 15, 20, and 25, and the experiment 1530 was performed under the same conditions as the atrazine and copper experiments described 1531 above.
- Photosynthetic efficiency was measured as maximum quantum yield of photosynthetic activity on days 0, 2, 5, 10, 15, 20 and 25 by pulse-amplitude modulated (PAM) fluorometry (MAXI Imaging PAM fluorometer, Walz, Germany). Maximum quantum yield is determined as the ratio of variable and maximum fluorescence ( $F_v/F_m$ ). Maximum fluorescence  $F_m$  is the maximum fluorescence emission level after a dark adaptation of 20 minutes, measured with a saturating pulse of light (emission peak at 450 nm, 2700 photons m<sup>-2</sup>s<sup>-1</sup>, 800 ms). Variable fluorescence  $F_v$  is calculated from the difference between initial fluorescence ( $F_0$ ) and maximum fluorescence ( $F_v = F_m - F_0$ ).
- 1539 EPS were measured by spectrophotometry at the end of the experiments after 25 days. Three 1540 replicates of 10 ml suspended diatom culture per treatment were centrifuged for 15 min at 15 °C 1541 and 3500xg. The supernatant yielded the soluble EPS fraction, which was left to precipitate 1542 overnight in 30 ml cold ethanol (98%), and subsequently centrifuged for 15 min at 15 °C and 1543 3500g. The pellet was dried under a flow of nitrogen and resuspended in 2 ml of 1.5% NaCl. 1544 Samples of 200 µl of this suspension per replicate were used for the EPS analysis. EPS was 1545 measured according to a modified version of the phenol/ $H_2SO_4$  assay (Dubois et al. 1956). In 24-1546 well-plates, 1 ml concentrated  $H_2SO_4$  and 200 µl phenol (5%, w / v in distilled water) were added 1547 to 200  $\mu$ l sample. The mixture was shaken, incubated for 20 min and the absorbance measured at 1548 488 nm using a Victor<sup>3</sup> multilabel reader (Perkin-Elmer). EPS concentrations were quantified with

a function obtained from a linear regression of known concentrations of glucose standards
(ranging from 5 to 400 μg/ml).

1551 Fatty acid profiles in 2 replicates per treatment were analyzed after 25 days. A suspension of 10 1552 ml suspended diatom culture was centrifuged for 20 minutes at 4 °C and 2700g. The supernatant 1553 was removed and the pellet was stored at -80 °C until fatty acid extraction. Hydrolysis of total lipid 1554 extracts of duplicate diatom cell suspensions per treatment and methylation to fatty acid methyl 1555 esters (FAME) was achieved by a modified one-step derivatisation method (Abdulkadir and 1556 Tsuchiya 2008; De Troch et al. 2012a). The boron trifluoride-methanol reagent was replaced by 1557 3.5 ml of a 2.5% H<sub>2</sub>SO<sub>4</sub>-methanol solution per sample since BF3-methanol can cause artefacts or 1558 loss of PUFA (Eder 1995). The fatty acid methylnonadecanoate C19:0 (Fluka 74208) was added 1559 as an internal standard for the quantification (see further). Samples were centrifuged and vacuum 1560 dried. The FAME thus obtained were analysed using a gas chromatograph (HP 6890N) coupled to 1561 a mass spectrometer (HP 5973). The samples were run in splitless mode, with a 1  $\mu$ L injection per 1562 run, at an injector temperature of 250°C using a HP88 column (Agilent J&W; Agilent Co., USA). The 1563 oven temperature was programmed at 50°C for 2 min, followed by a ramp at 25°C min<sup>-1</sup> to 175°C 1564 and then a final ramp at 2°C min<sup>-1</sup> to 230°C with a 4 min hold. The FAME were identified by 1565 comparison with the retention times and mass spectra of authentic standards and mass spectral 1566 libraries, and analysed with the software MSD ChemStation (Agilent Technologies). The 1567 quantification function of each individual FAME was obtained by linear regression of the 1568 chromatographic peak areas and corresponding known concentrations of the external standards 1569 (Supelco # 47885, Sigma-Aldrich Inc., USA) ranging from 5 to 250 ng/ml).

All traits and functional contribution measurements were normalized to cell density and areindicated in Addendum II Table S1.

1572 **Data analysis.** The numerical stress response was quantified as EC50, i.e. the effective copper and 1573 atrazine concentration inducing a 50% loss of each species' maximum cell density (carrying 1574 capacity). The functional stress response was quantified as the EC50 of photosynthetic efficiency, 1575 EPS and fatty acids production, representing the effective concentration of copper and atrazine 1576 inducing a 50% reduction in these processes. EC50 values as a sensitivity endpoint represent the 1577 baseline for pesticide risk assessment in Europe (Van den Brink et al. 2011) and were calculated 1578 using a three-parameter log-logistic model (Equation 1) and a three-parameter Weibull model in 1579 two different parameterisations (Equations 2 and 3) as described in Ritz and Streibig (2005).

1580  $Y = d / (1 + \exp\{b[\log x - \log e]\})$  [1]

1581 
$$Y = d(1 - \exp[b(\log x - \log e)])$$
 [2]

1582  $Y = d(\exp\{-\exp[b(\log x - e)]\})$ 

1583 *Y* represents the response variable (cell density, photosynthetic efficiency, EPS or fatty acids per 1584 cell), *b* the relative slope of the curve, *d* the upper limit, *e* the inflection point, and *x* the 1585 concentration of atrazine or copper.

[3]

To account for the hormetic growth and EPS production of several species under stress, doseresponse models that can describe hormesis were fitted as described by Brain and Cousens (1989, Equation 4) and Cedergreen et al. (2005, Equation 5).

1589 
$$Y = c + (d - c + fx)/(1 + \exp\{b[\log x - \log e]\})$$
 [4]  
1590  $Y = (d + f \exp(-1 / x^{\alpha})) / (1 + \exp(b(\log x - \log e)))$  [5]

1591 *c* is the lower and *b* the upper horizontal asymptote and *f* is the hormesis effect. Model 5 was fitted 1592 with  $\alpha = 1$ , 0.5 and 0.25, respectively. Model fit for every dose-response model was evaluated 1593 visually as well as by likelihood ratio testing, lack-of-fit test and Akaike information criterion 1594 (AIC), and the best-fitting model was used to calculate EC50 values.

Trait-sensitivity relations were analyzed with generalized least squares models (Equation 6), with numerical and functional response to copper and atrazine as response variable and traits as predictors (biovolume, surface to volume ratio, length, cholorophyll a content, mixotrophy at 0.1 and 0.6 mM Glucose, TN and TOC).

1599 
$$S = a + \sum (b_i \cdot T_i)$$
 [6]

1600 T<sub>i</sub> represents the traits, S the numerical and functional response (EC50) in terms of respectively 1601 cell density, and photosynthetic efficiency, EPS and fatty acid production, a the intercept and b<sub>i</sub> 1602 the slope (i.e. the effect of trait i on stress response). Generalized least squares models were fitted 1603 to all uncorrelated traits (correlation factor < 0.5) as predictors, correlated traits were separately 1604 included as predictors in additional model fits. In all models, predictors were removed in a 1605 stepwise manner, model fit was assessed at every step (AIC and likelihood ratio testing), and the 1606 best-fitting model was selected for the trait-sensitivity analyses. When a trait significantly 1607 predicted numerical or functional stress response, we tested in additional analyses if the 1608 respective trait also predicted stress response when species with outlier trait values were 1609 omitted.

1610 The relation between the functional contribution in unstressed conditions and the numerical as 1611 well as functional stress response was analyzed by fitting generalized least squares models to 1612 functional contribution per cell as response variable and numerical as well as functional stress1613 response as predictors (Equation 7).

[7]

[8]

1614  $F = a + b \cdot S_D + c \cdot S_F$ 

F is the functional contribution (photosynthetic efficiency, fatty acid and EPS production per cell under control conditions),  $S_D$  is the numerical and  $S_F$  the functional stress response, a is the intercept, b and c represent the slopes, i.e. the relation of functional contribution to numerical and functional stress response. If  $S_D$  and  $S_F$  were correlated (correlation factor > 0.5), models were fitted separately for both predictors.

1620 The relation of numerical and functional stress response was assessed by fitting generalized least 1621 squares models to numerical response as predictor and the response of the respective function as 1622 response variable (Equation 8).

1623 
$$S_F = a + b \cdot S_D$$

S<sub>F</sub> is the functional stress response in terms of photosynthetic efficiency, EPS and fatty acid
 production respectively, a is the intercept, S<sub>D</sub> the numerical stress response in terms of cell
 density, b represents the slope, i.e. the relation of numerical and functional stress response.

- 1627 For all generalized least squares models, normality and homogeneity of model residuals were 1628 evaluated using quantile-quantile plots and by plotting residuals versus explanatory variables, 1629 respectively. If homogeneity was violated, the model was refitted using an exponential variance 1630 structure allowing residuals to change separately with each predictor X (var( $\varepsilon$ ) =  $\sigma 2 \cdot e^{2 \cdot \delta \cdot X}$ ), the 1631 best-fitting model (lowest AIC) was selected, and homogeneity was tested again. By means of 1632 likelihood ratio testing, the significance of this structure was tested (alpha = 0.05). All calculations 1633 were performed in R 2.10.1, using RStudio and the packages drc and nlme (Ritz and Streibig 2005, 1634 Pinheiro et al. 2015, R Development Core Team 2016).
- 1635

#### 1636 **3.3. Results**

The numerical response of diatoms to copper was strongly related to biovolume, with larger species being more tolerant to copper (p<0.0001, Fig. 3.1a, Addendum II Table S3). Other traits that were correlated with biovolume (surface to volume ratio, length, chlorophyll a content) also predicted the numerical response to copper in separate model fits (Addendum II Table S4). In contrast, the numerical response to atrazine was not related to any morphological trait, but to the capacity for mixotrophic growth (p=0.0003, Fig. 3.1b, Addendum II Table S3). Diatom species which could increase their density in response to glucose turned out to be least affected by atrazine (see mixotrophic growth response and atrazine sensitivity per species in Addendum II
Tables S1 and S5). Biovolume and mixotrophy still predicted the numerical response to copper
and atrazine in additional analyses where the largest respectively most mixotrophic species
(*Entomoneis* sp. and *Cylindrotheca closterium*, data points in upper right quadrant of Fig. 3.1a and
3.1b) were omitted (Addendum II Table S4).

1649 The functional response to copper was predicted by biovolume for all three ecosystem processes 1650 measured, with larger species being least affected in their photosynthetic efficiency, fatty acid 1651 content and EPS production (all p<0.0001, Fig. 3.1c, Addendum II Table S3). The functional 1652 response to atrazine in terms of photosynthetic efficiency was also positively related to biovolume 1653 (p<0.0001, Fig. 3.1d, Addendum II Table S3). The same outcome was obtained when the largest 1654 species (Entomoneis sp.) was omitted from the analysis (Addendum II Table S4). In separate 1655 model fits, other morphological traits that were correlated with biovolume (surface to volume 1656 ratio, length) also predicted the response of photosynthetic efficiency, EPS and fatty production 1657 to copper as well as the response of photosynthetic efficiency to atrazine (Addendum II Table S4). 1658 None of the measured traits predicted the functional response to atrazine in terms fatty acid and 1659 EPS production (Fig. 3.1d, Addendum II Table S3), if outlier data were omitted from the analysis 1660 (Addendum II Table S4). EC50s for all diatom strains are indicated in Addendum II Table S5, and 1661 the growth curves for every strain and stressor are shown in Addendum II Fig. S2. The diatoms' 1662 morphological traits were not altered during exposure to atrazine and copper, with the exception 1663 of Astartiella bahusiensis, which increased its cell size when exposed to copper (Addendum II 1664 Table S8 and Fig. S1).

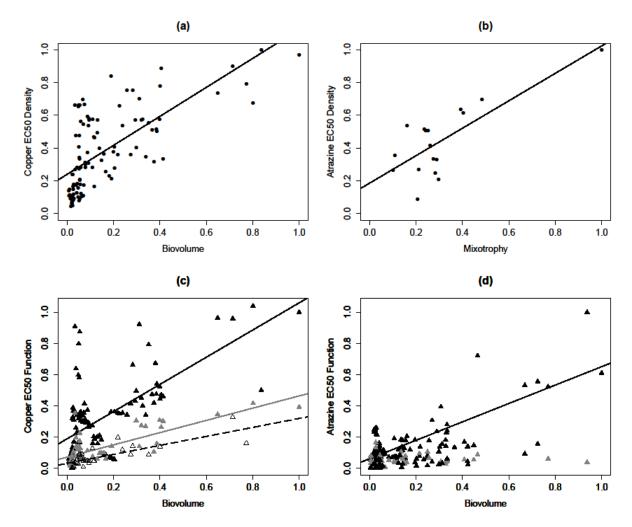


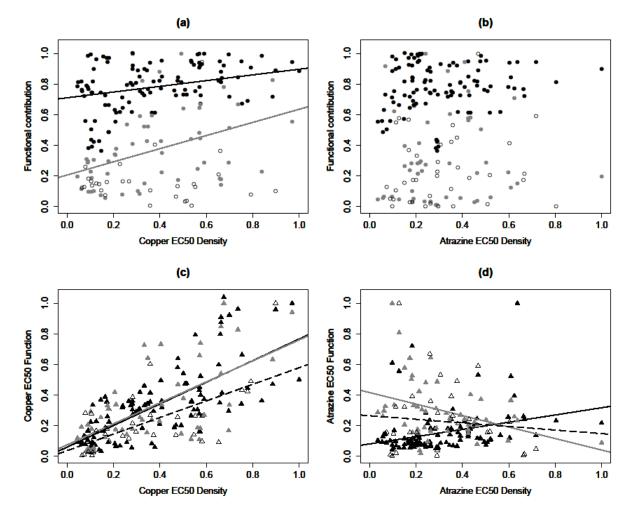


Fig. 3.1: Diatom numerical and functional response to copper and atrazine as a function of biovolume and 1666 1667 mixotrophy. Panel a) shows numerical response to copper (EC50 cell density) as a function of biovolume. 1668 Panel b) shows numerical response to atrazine as a function of mixotrophy. Regression lines show 1669 significant relations between biovolume or mixotrophy and numerical response to copper and atrazine. 1670 Panels c) and d) show functional response (EC50 of photosynthetic efficiency, fatty acids and EPS production) to copper and atrazine as a function of biovolume. Regression lines show significant relations 1671 1672 between biovolume and functional response: photosynthetic efficiency (black triangles, black lines), fatty 1673 acids (white triangles, dashed line), and EPS production (grey triangles, grey line). Biovolume, mixotrophy 1674 and EC50 values are indicated normalized to the highest respective value recorded.

1675

The functional contributions of diatoms in unstressed conditions were not correlated to their numerical or functional response to atrazine for any of the three processes (Fig. 3.2b, Addendum II Table S6). Diatom numerical and functional response to copper was positively related to photosynthetic efficiency (p=0.0008 for numerical and p=0.0006 for functional response) as well as EPS production (p=0.001 for numerical and p=0.009 for functional response), with productive species being more copper-resistant, but not fatty acid content (Fig. 3.2a, Addendum II Table S6).

1682



1683

1684 Fig. 3.2: Diatom functional contribution and functional stress response as a function of numerical response 1685 to copper and atrazine. Panels a) and b) show functional contribution in unstressed conditions as a function 1686 of numerical response (EC50 cell density) to copper and atrazine. Significant relations are indicated by 1687 regression lines. Functional contributions are photosynthetic efficiency (black dots, black line), fatty acids 1688 (white dots) and EPS production (grey dots, grey line). Panels c) and d) show the relation between 1689 numerical response and functional response (EC50 of photosynthetic efficiency, fatty acids and EPS 1690 production) to copper and atrazine. Significant relations are indicated by regression lines. Functional 1691 response is indicated in terms of photosynthetic efficiency (black triangles, black line), fatty acids (white 1692 triangles, dashed line) and EPS production (grey triangles, grey line). All functional contribution and EC50 1693 values are indicated normalized to the highest respective value recorded.

1694

1695 The numerical and functional stress response were related, this relationship however differed 1696 between copper and atrazine. For copper, species which were least affected in their density were 1697 also the most tolerant in their functional contribution. Numerical and functional response to 1698 copper were positively related for all three processes (p < 0.0001 for photosynthetic efficiency and 1699 EPS production, p=0.0015 for fatty acid content, Fig. 3.2c, Addendum II Table S7). A similar 1700 positive relation was observed between the numerical response and the response of 1701 photosynthetic efficiency to atrazine (p=0.003, Fig. 3.2d, Addendum II Table S7). Conversely, the 1702 numerical and functional response to atrazine in terms of EPS and fatty acid production were 1703 negatively related (p=0.0015 and p=0.03, Fig. 3.2d, Addendum II Table S7). These negative 1704 relations between the numerical and functional response to atrazine were largely driven by 1705 species whose density was least affected by atrazine (e.g. *Cylindrotheca closterium, Gyrosigma* sp. 1706 2, Cylindrotheca fusiformis, Biremis ambigua), but which were amongst the most atrazine-sensitive 1707 ones concerning their functional contribution (Addendum II Table S5). The functional response 1708 to copper and atrazine stress was thus not only predicted by traits, but was also related to the 1709 species' capacity to maintain their density under both types of stress: the numerical and functional 1710 response to atrazine were not driven by the same set of traits, and atrazine-resistant species in 1711 terms of density were amongst the most sensitive in their EPS and fatty acid output. The numerical 1712 and functional response to copper were driven by the same morphological traits, and numerically 1713 resistant species were least affected in their functional contribution.

1714

### 1715 **3.4. Discussion**

1716 The outcome of our experiments demonstrates that the numerical and functional response to two 1717 common chemical stressors can be predicted from traits, but also that these stressors are likely to 1718 have a differential impact on ecosystem functioning as the relationship between response and 1719 effect traits is different depending on stressor and functional process. When exposed to copper, 1720 response and effect traits were correlated and the numerically tolerant species were least affected 1721 in their functioning, suggesting that under metal stress the first species to be numerically 1722 decimated will probably have little effect on overall ecosystem functioning. In contrast, 1723 mixotrophic growth can help to maintain the density of certain species in response to 1724 photoinhibiting stressors such as atrazine, but not their contribution to particular functional 1725 processes, which would cause the functional consequences of stress exposure to be more 1726 important than estimated from mere species densities.

1727 Trait-based approaches have made a considerable breakthrough in explaining community 1728 structure along gradients of environmental stress (Baird and Van den Brink 2007, Litchman et al. 1729 2007, Fontana et al. 2014). Yet, effect traits leading to increased or decreased ecosystem 1730 functioning are rarely linked the response traits needed to maintain density under stress 1731 (Hillebrand and Matthiessen 2009, but see e.g. Pomati and Nizzetto 2013, Heuner et al. 2015). This 1732 represents a major shortcoming for predicting ecosystem functioning in stressed environments, 1733 given that stress can affect ecosystem functioning by altering both the density and the functional 1734 output of the concerned species (Fox 2006). For this challenge, morphological traits are expected 1735 to be suitable tools, since they are relatively simple to measure, capture most of the functional 1736 properties of microalgae, and can play an important role in algae-pollutant interactions (Kruk et 1737 al. 2010).

1738 In the case of copper, the expectations of morphological traits as predictors of any type of stress 1739 response were fulfilled as biovolume, surface to volume ratio and cell length all predicted the 1740 response of diatom density, photosynthetic efficiency, fatty acid and EPS production to copper. 1741 The case of atrazine however demonstrated that morphological traits alone can not generally 1742 predict numerical and functional response. As common chemical stressors in coastal and 1743 estuarine environments, the effects of atrazine and other photoinhibiting herbicides on primary 1744 producers in these ecosystems have been extensively analysed, but mainly on the density and not 1745 on the functions sustained by the species concerned, and in relation to taxonomy rather than 1746 species traits. Mixotrophy is common in benthic diatoms (Tuchman et al. 2006), and the few 1747 approaches which related the effects of photoinhibiting herbicides to mixotrophy suggest the 1748 numerical response of diatoms to these stressors can be modulated by their trophic mode 1749 (Debenest et al. 2009; Larras et al. 2014). Here, mixotrophy predicted the numerical response of 1750 diatoms, but not their contribution to any of the three functional processes under atrazine. 1751 Estimating the effects of photoinhibiting stress on ecosystem functions sustained by primary 1752 producers is therefore less straightforward than under metal stress, since the numerical and 1753 functional response to photoinhibiting stress are not driven by the same traits and mixotrophy 1754 cannot be quantified as easily as morphological traits. Nevertheless, the capacity for mixotrophic 1755 growth can be obtained from the literature for many taxa (Hellebust and Lewin 1977, Tuchman 1756 et al. 2006). Our study indeed corroborates literature reports for Cylindrotheca closterium 1757 (Vanelslander et al. 2009), which was also the most mixotrophic species in our experiments.

1758 Relating traits to stress response should be done by linking traits to groups of stressors with 1759 similar modes of action, to generalize the trait-sensitivity relation for distinct types of stressors 1760 (Baird and Van den Brink 2007). Like copper, the toxicity of other heavy metals to primary 1761 producers lies in the formation of reactive oxygen species (ROS), which can damage cell 1762 membranes, plastids and other intracellular structures (Masmoudi et al. 2013). Metal uptake is 1763 commonly related to cell size, with smaller species having a larger surface to volume ratio and 1764 thus being more prone to be adversely affected by metal pollution (Miao et al. 2005, Quigg et al. 1765 2006). Not only morphological traits, but also protective mechanisms such as the production of 1766 metallothioneins equally play an important role in mediating metal toxicity in various organisms 1767 (Cobbett and Goldsbrough 2002). Nevertheless, morphological traits appear as a promising, 1768 general tool to estimate both the numerical and functional response of microalgae to metal 1769 pollution.

1770 In contrast to metals, triazine herbicides are photoinhibiting stressors, which disrupt
1771 photosynthesis rather than causing physical damage to cell structures. In this respect, the effects
1772 of triazine herbicides resemble those of other types of photoinhibiting stress in benthic intertidal

1773 habitats, caused by extreme light conditions due to tidal emersion in combination with 1774 unpredictably changing weather conditions (Barnett et al. 2015). Within growth forms 1775 constraints (motile vs non-motile), diatoms can adapt to the excess light stress by behavioural 1776 (e.g. vertical migration in motile species) and/or physiological photoprotective mechanisms such 1777 as non-photochemical quenching (McQuoid and Hobson 1996, McLachlan et al. 2009, Barnett et 1778 al. 2015). However, in contrast with mixotrophic capacity, none of these adaptations would 1779 probably be efficient against herbicide-induced blocking of PS II. Phylogenetic indices integrate 1780 over other traits that can potentially mediate algal resistance to PS II inhibitors (e.g. absence or 1781 presence of a raphe structure, Larras et al. 2014) and are therefore, alongside the capacity for 1782 mixotrophic growth, suggested as promising predictors of microalgae response to herbicide 1783 stress (Larras et al. 2014).

1784 The variability of traits and the multitude of potential sensitivity endpoints should be considered 1785 when using trait-based approaches to predict stress response (Albert et al. 2010, Rubach et al. 1786 2012). We observed little changes in morphological traits under stress exposure, but our 1787 experiments lasted only a limited number of vegetative growth cycles and hence negligible size 1788 reduction (inherent to the diatom life cycle, cf. Chepurnov et al. 2008). Diatoms can shift their 1789 trophic mode with almost immediate effect (Lewin and Hellebust 1975), whereas significant 1790 changes in morphology occur over longer time spans which makes trait variability more likely in 1791 long-term experiments and in scenarios of constant stress exposure. Also, in toxicology, sensitivity 1792 is defined primarily as the median of a tolerance distribution (the 'EC50') along an environmental 1793 gradient (Baird et al. 1996). Conversely, in ecology, sensitivity is often defined as the variance of 1794 the tolerance distribution, irrespective of its median, with respect to an environmental gradient 1795 (Baird et al. 1996). While less straightforwardly linked to non-morphological traits than, for 1796 example, toxicokinetic measurements (e.g. uptake or elimination rate constants, Rubach et al. 1797 2012), EC50 values were selected as sensitivity endpoints because of their widespread use in 1798 trait-sensitivity research (Baird and Van den Brink 2007; Van den Brink et al. 2011; Larras et al. 1799 2014). However, our analysis could also be performed based on other sensitivity endpoints that 1800 better capture the variance of the tolerance distribution and relate to a wider variety of traits 1801 (Rubach et al. 2012).

To evaluate the effects of different stressor types on ecosystem functioning, we need to contrast and disentangle the response traits driving the species densities and the effect traits driving the contribution to ecosystem functioning under stress. Mixotrophy as a response trait predicted diatom densities, but not functioning under atrazine exposure, and those species which could best maintain their cell densities and photosynthetic efficiency under atrazine were most sensitive in their fatty acid content and EPS production. These contrasting effects of atrazine on density and 1808 functioning imply that at herbicide stress levels where species densities are not yet impacted, 1809 sediment stabilization and energy flux to higher trophic levels would already be affected, with 1810 concomitant knock-on effects on other sediment-associated organisms and the food web in 1811 intertidal systems (Müller-Navarra et al. 2000, Tolhurst et al. 2003, De Troch et al. 2012a).

1812 For metal pollution, the functional consequences seem at a first sight smaller: response and effect 1813 traits were positively correlated, meaning that species that were little affected in their density also 1814 performed best at maintaining their contribution to the functional processes. Furthermore, 1815 copper mostly reduced the densities of those species that contributed least to EPS and fatty acids 1816 production. Since the same traits that predicted the numerical and functional response to copper 1817 in this study also explain tolerance to pollution by other metals (Miao et al. 2005, Lahive et al. 1818 2011), large, metal-resistant species could potentially sustain ecosystem functioning under metal 1819 pollution. However, the real threat for species loss and ecosystem functioning is from the presence 1820 of multiple stressors that primary producers are subjected to in intertidal environments 1821 (Underwood et al. 1998, Hillebrand et al. 2000, McLachlan et al. 2009, Souffreau et al. 2010). Cell 1822 size is a master trait that shapes the performance of microalgae along environmental gradients 1823 (Litchman and Klausmeier 2008, Pomati and Nizzetto 2013, Pomati et al. 2013), but large size is 1824 correlated with other extinction-prone traits such as lower nutrient acquisition, slower growth 1825 and longer generation times (Litchman and Klausmeier 2008, Pomati et al. 2013, Litchman et al. 1826 2015). If simultaneously with metal pollution diatoms are subjected to other types of stress that 1827 favour traits such as small cell size or short growth cycles, the competitive advantage and the 1828 capacity of large-bodied species to sustain ecosystem functioning will be limited.

1829 Lastly, we should consider the relation between numerical and functional stress response not only 1830 from an ecosystem, but also from a diatom perspective. The processes performed by diatom 1831 biofilms are essential contributions to ecosystem functioning (sediment stabilization, primary 1832 production, energy transfer to higher trophic levels). But for the diatoms themselves, a high 1833 photosynthetic efficiency will be beneficial under most types of growth-limiting stress, whereas 1834 for example producing large amounts of fatty acids will not be advantageous in the case of 1835 herbicide stress. Oxidative stress resulting from metal exposure on the other hand results in the 1836 damage of fatty acid rich structures such as cell membranes due to lipid peroxidation (Li et al. 1837 2006), and a high fatty acid production could help diatoms to better tolerate the damage to cellular 1838 structures caused by heavy metal stress. Similarly, producing EPS can also protect diatoms form 1839 metal stress, through the establishment of an organic carbon matrix that protects biofilms by 1840 chelating metal ions (Pistocchi et al. 1997). Generally, stressed organisms will allocate most 1841 resources to those processes that favour survival and growth under stress and we should consider

the interconnection of numerical and functional stress response, when interpreting theirdivergent relation under different types of stress.

1844 In conclusion we infer two general statements from our results. First, the numerical and functional response of primary producers to two of the most common chemical stressors in coastal 1845 1846 environments, herbicide and metal pollution, is predictable based on the species' morphological 1847 traits and trophic mode. Second, the impacts of stress exposure on ecosystem functioning are 1848 underpinned by the relation of the numerical and functional stress response and the functional 1849 importance of those species whose density is reduced first. Linking response and effect traits 1850 driving the numerical and functional stress response will reveal the capacity of tolerant species to 1851 maintain their contribution to ecosystem functioning, and should be considered when estimating 1852 ecosystem functioning under global change.

1853

# 1854 Chapter 4: Facilitation and tolerance explain the diversity effect on 1855 functioning in stressed diatom communities

1856

## 1857 Abstract

The relation between biodiversity and ecosystem functioning depends on complementarity and
dominance effects, but the ecological processes driving complementarity and dominance under
anthropogenic stress remain largely unknown.

We exposed two benthic diatom communities to metal and pesticide stress (copper and atrazine), and measured complementarity and dominance effects on biomass yield under stressed and unstressed conditions. Next, we analysed if complementarity under stress was 'two-way' (affecting species independent of their biomass yield) or 'one-way' (differentially affecting highand low-yield species). Last, we tested if the production of extracellular polymers as a facilitative mechanism as well as the species' stress tolerance and capacity for mixotrophic growth were ecological processes that explained complementarity and dominance under stress.

The net biodiversity effect on diatom biomass yield was consistently positive, and was mainly caused by two-way complementarity, which was higher in diatom communities stressed by copper (both communities) and atrazine (one community) than in unstressed communities. Stimulation by stress of the diatom's extracellular polymer production predicted, in part, these increases of complementarity under stress.

1873 Complementarity in communities stressed by the atrazine, a photosynthesis inhibitor, was driven 1874 by the presence of mixotrophic species, which can grow on organic substances. Under copper 1875 stress, complementarity in one community was driven by the species which were least tolerant to 1876 the metal in monoculture. These species were characterised by low biomass yields, resulting in a 1877 negative one-way complementarity, reducing the net biodiversity effect under metal stress.

*Synthesis.* Biodiversity effects on diatom biomass were caused by two-way complementarity effects, which increased under stress and were driven by extracellular polymer production, a 'sleeping' facilitative mechanism stimulated by stress. The degree to which diatom species benefited from complementarity depended on their trophic mode and stress tolerance. When complementarity favoured the growth of sensitive but low-yield species, the net biodiversity effect on diatom biomass yield decreased under stress.

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1885

#### 1886 **4.1. Introduction**

Understanding the influence of biodiversity on ecosystem functioning is a key challenge, as global
change is causing unprecedented rates of biodiversity change (Pereira et al. 2010, Dornelas et al.
2014, Gonzalez et al. 2016). A large amount of empirical research has documented in most cases
positive effects of biodiversity on ecosystem processes (most commonly biomass yield, Balvanera
et al. 2006, Cardinale et al. 2011, Naeem et al. 2012), with results being largely consistent across
trophic levels and ecosystem types (Cardinale et al. 2006, Griffin et al. 2013).

1893 Anthropogenic stressors are a main driver of biodiversity change (Wilcove and Master 2005, Malaj 1894 et al. 2014), and can substantially affect biodiversity-ecosystem functioning relationships (Steudel 1895 et al. 2012, Baert et al. 2016). An increasing number of studies have tested the biodiversity effect 1896 on ecosystem functioning under stressful conditions, but these have yielded contrasting results 1897 (clearly positive effects in Mulder et al. 2001, Ives and Cardinale 2004, Solan et al. 2004, Zavaleta 1898 and Hulvey 2004, Larsen et al. 2005, McMahon et al. 2012, Steudel et al. 2012, Mensens et al. 2015, 1899 Baert et al. 2016 and neutral or even negative diversity effects in Jonsson et al. 2002, Petchey et 1900 al. 2002, Smith and Knapp 2003, Jiang 2007, Jiang et al. 2008, Roger et al. 2012). This variability 1901 of results calls for more mechanistic studies on how stress affects diversity effects on functioning, 1902 relating ecological processes to stress-induced alterations of biodiversity effects on function. 1903 (Naeem 2008, Solan et al. 2009, O'Connor et al. 2015, Baert et al. 2016).

1904 Dominance (dominance of communities by high- or low-yield species) and complementarity 1905 (better performances of species in community due to mechanisms such as resource partitioning 1906 or facilitation) are the main diversity effects which cause communities to perform differently than 1907 expected from monocultures of the constituent species (Loreau and Hector 2001, Fox 2005). 1908 Dominance and complementarity mostly operate simultaneously (Loreau et al. 2001). 1909 Consequently, which effect drives the diversity-functioning relation and how both effects are 1910 affected by anthropogenic stress is actively debated (Cardinale et al. 2007, Fargione et al. 2007, 1911 Jiang et al. 2008, Loreau 2010, Wang et al. 2013, Baert et al. 2016, Hodapp et al. 2016). 1912 Complementarity and dominance effects can drive ecosystem functioning under stressful 1913 conditions through stress-induced increases of facilitation (Wang et al. 2013) and when 1914 communities are dominated by stress-tolerant species with above-average contribution to 1915 functioning (Baert et al. 2016), respectively. Even though still scarce at the moment, stressor-1916 induced changes in dominance and complementarity are increasingly tested for (Wang et al. 2013, 1917 Baert et al. 2016), using analytical methods based on the covariance between species performance 1918 in a community and in monoculture (Loreau and Hector 2001, Fox 2005).

1919 On itself, quantifying complementarity and dominance does not offer direct mechanistic insight 1920 into the ecological processes driving stress effects on biodiversity-ecosystem functioning 1921 relationships. That is because complementarity and dominance (sensu Fox 2005 or Loreau and 1922 Hector 2001, who refer to complementarity and selection) are statistical effects that can result 1923 from multiple ecological processes (Cardinale et al. 2011, Tilman et al. 2014). One such process 1924 is facilitation, which can drive complementarity in legumes, insects and microalgae (Cardinale et 1925 al. 2002, Temperton et al. 2007, Vanelslander et al. 2009). Further tests of facilitative mechanisms 1926 as drivers of complementarity are however lacking and there is no direct evidence attributing 1927 changes of complementarity effects under stress to changing facilitative mechanisms.

1928 Here, we quantify complementarity and dominance effects on the biomass yield of two benthic 1929 diatom communities under chemical stress, and compare these with effects determined in 1930 unstressed controls. We use benthic diatom communities as an experimental model because they 1931 are the main primary producers in many intertidal, soft-sediment habitats, and diatom biomass 1932 production is crucial as it represents the basis of the food web in these systems (Montagna et al. 1995, Underwood and Kromkamp 1999, Tang and Kristensen 2007). We use copper and the 1933 1934 herbicide atrazine as model stressors, the former being a common pollutant in coastal ecosystems 1935 due to urban and industrial runoff (Kennish 1996, Murray-Gulde et al. 2002), and the latter 1936 representing one of the most-used agricultural herbicides worldwide (Graymore et al. 2001, 1937 Chirnside et al. 2007).

1938 We then test if complementarity under stress was 'two-way' (affecting diatom species 1939 comparably) or 'one-way' (differentially affecting species with high or low biomass yields). 1940 Benthic diatoms produce large amounts of extracellular polymeric substances (EPS, Decho 1990, 1941 Underwood and Paterson 2003). These organic exudates have been linked to metal tolerance in 1942 diatom monocultures (Pistocchi et al. 1997, Levy et al. 2007) and can cause facilitation in diatom 1943 communities containing mixotrophic species (Vanelslander et al. 2009). We therefore analyse if 1944 the release of EPS could predict complementarity in stressed diatom communities, and test if 1945 stressor-induced changes in complementarity were related to the species' stress tolerance in 1946 monoculture and the presence of mixotrophic species.

1947

#### 1948 **4.2. Methods**

Experimental organisms and culture conditions. The two diatom communities (hereafter
community A and B) were each composed of six benthic diatom species (Addendum III Table S1)
collected in the Westerschelde estuary (The Netherlands, 51°21'N, 3°43'E) and the Veerse meer
(The Netherlands, 51°32'N, 3°47'E), where they represent the dominant component of the

microphytobenthos (Forster et al. 2006). Monoclonal cultures of each strain were obtained and
preserved in the culture collection of the Protistology & Aquatic Ecology Research Group (UGent,
http://bccm.belspo.be/). Diatoms were maintained in a climate room at 15±1 °C, a light/dark
cycle of 12h / 12h and an illumination of 90 µmol photons m<sup>-2</sup>s<sup>-1</sup>, in culture medium consisting of
filtered and autoclaved natural seawater (salinity 32±1) enriched with f/2 nutrients (Guillard
1958 1975).

1959 Atrazine and Copper experiments. Diatoms were exposed to 0 (control), 200 and 500 µg/l 1960 (hereafter 'low' and 'high') atrazine and copper respectively. In a previous experiment, these 1961 concentrations resulted on average in a 50% and 90% reduction of diatom monoculture yield 1962 (Chapter 3). Atrazine treatments were prepared from a stock solution obtained by dissolving 50 1963 mg technical atrazine (2-chloro-4-ethylamino-6-isopropylamino-s-triazine, 99.8% pure, Sigma-1964 Aldrich Chemie Gmbh, Munich, Germany) in 10 ml acetone as a carrier to increase the solubility 1965 of atrazine, with a maximum final volume of 0.01% acetone in the treatments. An acetone control 1966 treatment of 0.01% acetone was included and compared to an acetone-free control to test for 1967 carrier effects. All atrazine treatments were compared to the acetone control. Copper (as a 1968 Cu[II]Cl<sub>2</sub> solution, analytical grade; VWR International) was spiked directly into the culture 1969 medium before exposure of the diatoms. F/2 culture medium for the copper experiments was 1970 prepared without EDTA, to avoid complexation of free copper ions (Pistocchi et al. 1997). The 1971 resulting atrazine and copper concentrations were determined by respectively GC-MS and 1972 graphite furnace AAS analyses (Addendum III Table S2). All experiments were run in tissue 1973 culture flasks (Greiner BioOne, CELLSTAR® TC, 175 cm<sup>2</sup> growth surface), with nine replicates per 1974 treatment (three replicates in the monoculture treatments). Each microcosm was inoculated with 1975 a total cell density of 5000 diatom cells/ml (5000 cells/ml for each species in monoculture, 1976 between 800-850 cells/ml per species in communities). Microcosms were inoculated with cells 1977 from exponentially growing cultures and were incubated in a climate room at 15±1 °C, under a 1978 light / dark cycle of 12h / 12h at 90 µmol photons m<sup>-2</sup>s<sup>-1</sup>. Culture medium of all treatments was 1979 refreshed after eight and 15 days, and experiments were terminated after 25 days.

1980 **Diatom biomass.** Diatom biomass was quantified as biovolume, and was calculated from cell 1981 densities and mean cell biovolume per species, using measured linear dimensions and formulas 1982 representing the closest approximation of geometric shape for each genus Hillebrand et al. (1999). 1983 Cell densities were obtained at day 0, 2, 5, 10, 15, 20 and 25 of the experiment by magnifying and 1984 photographing (x100) an area of 0.66 mm<sup>2</sup> per microcosm, using an inverted Axiovert 135 Zeiss 1985 microscope (Carl Zeiss, Jena, Germany) and a connected digital camera (Canon PowerShot G11 1986 digital camera). Cell numbers per species were subsequently determined by identifying and 1987 counting cells on the computer (ImageJ cell counting software, Schneider et al. 2012) and converted into cell densities in cells/ml. All further analyses use the biomass on day 15 (lateexponential growth phase).

1990 **EPS production.** Diatom biofilms are characterized by a high production of EPS (Underwood and 1991 Paterson 2003), which represent a mix of organic compounds consisting mostly of 1992 polysaccharides, but also proteins and nucleic acids (Baines and Pace 1991). These organic 1993 compounds play an important role for sediment stabilization in muddy intertidal areas (Decho 1994 2000, Underwood and Paterson 2003). The release of organic compounds into the medium has 1995 also been identified as driver of complementarity effects in benthic diatoms (Vanelslander et al. 1996 2009). Furthermore, benthic diatoms can modulate their EPS production in response to abiotic 1997 stress, notably copper (Pistocchi et al. 1997, Levy et al. 2007). We therefore tested whether the 1998 EPS released by diatoms into the medium could predict potential changes of complementarity 1999 under chemical stress. EPS were measured by spectrophotometry after 15 days, in three 2000 replicates of 10 ml suspended diatom culture per treatment. The suspended diatom cultures were 2001 centrifuged for 15 min at 15 °C and 3500xg. The supernatant yielded the soluble EPS fraction, 2002 which was left to precipitate overnight in 30 ml cold ethanol (98%), and subsequently centrifuged 2003 for 15 min at 15 °C and 3500xg. The pellet was dried under a flow of nitrogen and resuspended in 2004 1 ml of 1.5% NaCl. Samples of 200 µl of this suspension per replicate were used for the EPS 2005 analysis. EPS were measured according to a modified version of the phenol/H<sub>2</sub>SO<sub>4</sub> assay by Dubois 2006 et al. (1956). In 24-well-plates, 200 µl phenol (5%, w / v in distilled water) and 1 ml concentrated 2007  $H_2SO_4$  were added to 200 µl sample. The mixture was shaken, incubated for 20 min and the 2008 absorbance measured at 488 nm using a Victor<sup>3</sup> multilabel reader (Perkin-Elmer). EPS 2009 concentrations were quantified with a function obtained from a linear regression of known 2010 concentrations of glucose standards (ranging from 5 to  $400 \mu g/ml$ ).

Data analysis. For every treatment, we partitioned the net biodiversity effect on diatom biomass
into dominance, two-way complementarity and one-way complementarity, using the method
developed by Loreau and Hector (2001) and later extended by Fox (2005, Equation 1).

$$\Delta Y = N \times \operatorname{cov} \left( M_i, \frac{RY_i}{RYT} - RY_{Ei} \right) + N \times (\overline{\Delta RY}) \times (\overline{M}) + N \times \operatorname{cov} \left( M_i, RY_i - \frac{RY_i}{RYT} \right)$$
[1]

2014  $\Delta Y$  is the net biodiversity effect on diatom biomass yield, i.e. the difference between the observed 2015 community yield and the community yield that is expected from the monoculture yields of the 2016 component species. M<sub>i</sub> is the mean monoculture yield of species *i*. *RY*<sub>i</sub> is the observed relative 2017 yield of species *i* with  $RY_i = \frac{O_i}{M_i}$ , where  $O_i$  is the observed biomass of species *i* in the community. 2018 *RYT* is the relative yield total with  $RYT = \sum_{i=1}^{N} RY_i$  where *N* is the total number of species in the 2019 community (six).  $\frac{RY_i}{RYT}$  is the observed frequency of species *i* in the community, i.e. its share of the 2020 relative yield total.  $\overline{M}$  is the average monoculture yield of all species.  $RY_{Ei} = \frac{1}{N}$  is the expected 2021 relative yield of species *i* in the community, with  $\sum_{i=1}^{N} RY_{Ei} = 1$ . The mean deviation in relative 2022 yield  $\overline{\Delta RY}$  was calculated as the mean difference in observed relative yield and expected relative 2023 yield  $\overline{\Delta RY} = \frac{1}{N} \times \sum_{i=1}^{N} (RY_i - RY_{Ei})$ .

The first term on the right-hand side of equation 1 is the 'dominance effect' (Fox 2005), which is the covariance between the species' monoculture yields and the difference between their observed and expected frequencies in community. The dominance effect is positive when species with high monoculture yields also reach high yields in community at the expense of low-yield species. It is negative if communities are dominated by low-yield species at the expense of highyield species. Mechanisms for this effect include interspecific competition for shared resources.

The second term is the 'two-way' or 'trait-independent' complementarity (TIC) effect (Loreau and Hector 2001, Fox 2005), which quantifies the extent to which the species' observed yields in community deviate from the expected yield, but in a way that is independent of the individual species' monoculture yields. Two-way complementarity is positive when growth in community rather than monoculture increases the yield of species, independent of their monoculture yield and not at the expense of other species. Mechanisms underlying this effect can be niche partitioning or facilitation.

2037 The third term is the 'one-way' or 'trait-dependent' complementarity (TDC) effect (Fox 2005), 2038 which is the covariance between the species' monoculture yield and the deviation of their 2039 observed relative yield from their observed frequency in community. One-way complementarity 2040 effects are observed when growth in community rather than monoculture increases the 2041 functioning of species with high or low monoculture yields, but not at the expense of other species. 2042 One-way complementarity can be positive if, for example, species with high monoculture yield are 2043 facilitated by species with low monoculture yield, but not vice-versa. One-way complementarity 2044 effects can become negative if low-yield species but not high-yield species benefit from facilitation 2045 when being grown in community.

To analyse if the net biodiversity effect, DOM, TIC and TDC changed under atrazine and copper stress, we used generalised least squares models with the biodiversity effects as response variables and the treatment type as categorical predictor (Equations 2-5).

2049	$\Delta Y = a + b \ge T$	[2]
2050	$DOM = a + b \times T$	[3]
2051	$TIC = a + b \times T$	[4]
2052	$TDC = a + b \times T$	[5]

 $\Delta Y$  is the net biodiversity effect on diatom biomass yield, DOM is the dominance effect, TIC and TDC are two-way and one-way complementarity, a is the intercept, T is the treatment type (control, low and high atrazine and copper) with the control set as reference level, b is the slope, i.e. the effect of the treatment type on each biodiversity effect.

2057 Next, we tested the role of EPS as a potential driver of complementarity in diatom communities
2058 by fitting generalized least squares models to two-way complementarity as response variable and
2059 diatom EPS production as predictor (Equation 6).

2060 TIC = 
$$a + b \times EPS$$
 [6]

TIC is two-way complementarity, a is the intercept, EPS the amount of EPS produced per unit
diatom biomass and b is the slope, i.e. the effect of EPS production on two-way complementarity.
Models were fitted separately for each diatom community and stressor.

In a last step, we tested if changes in one-way complementarity were driven by species with a low or high monoculture yield and/or stress tolerance benefiting most from being grown in community. We fitted generalized least squares models to the monoculture yield and stress tolerance of the diatom species as predictors, and the deviation of their observed relative yield from their observed frequency in community as response variable (Equation 7)

$$RY_{i,j} - \frac{RY_{i,j}}{RYT_j} = a + b \times M_{i,j} + c \times \frac{M_{i,j}}{M_{i,j=0}}$$
<sup>[7]</sup>

The fraction on the left-hand side of Equation 7 is the deviation of the observed relative yield from the observed frequency of species *i* when grown in community at stress level *j*. This term (hereafter relative performance) quantifies the extent to which each species was effected by oneway complementarity (see Equation 1).  $M_{i,j}$  is the monoculture yield of species *i* at the stress level  $j, \frac{M_{i,j}}{M_{i,j=0}}$  is the tolerance of species *i*, quantified is the ratio of the monoculture yield at stress level *j* and the monoculture yield in the control (*j*=0), a is the intercept, b and c are the slopes, i.e. the relation between the species' yield and tolerance in monoculture and the deviation of their
relative performance in community. If the monoculture yield and tolerance were correlated
(correlation factor > 0.5), models were fitted separately for both predictors. Models were fitted
separately for each diatom community and stressor.

2079 For all least squares model fits, normality and homogeneity of model residuals were inspected by 2080 evaluation of quantile-quantile plots and Shapiro-Wilk's test, and by Levene's test and plotting 2081 residuals versus explanatory variables respectively. If normality was violated (for model fits in 2082 equation 7 only), data were log-transformed. If homogeneity was violated, the model was refitted 2083 using a variance structure allowing different variances per treatment T ('weights = 2084 varIdent(form ~ 1 | T' in the R package nlme (Pinheiro et al. 2015) and homogeneity was 2085 evaluated again. By means of likelihood ratio testing, the significance of these structures was 2086 tested (alpha = 0.05). All calculations were performed in R 3.0.1. using RStudio (R Development 2087 Core Team 2016).

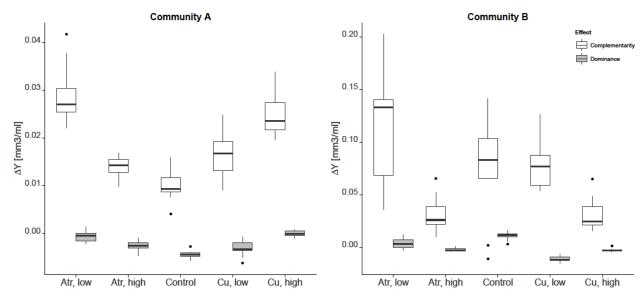
2088

### 2089 **4.3. Results**

Effects of stress on dominance and complementarity. The net biodiversity effect was positive in all treatments of both diatom communities, i.e. the biomass yield of diatoms in community was higher than that expected from the component species' monocultures (Addendum III Fig. S1). In both communities and across all treatments, complementarity was the main effect causing the yield of diatom communities to deviate from that expected from their monocultures (Fig. 4.1).

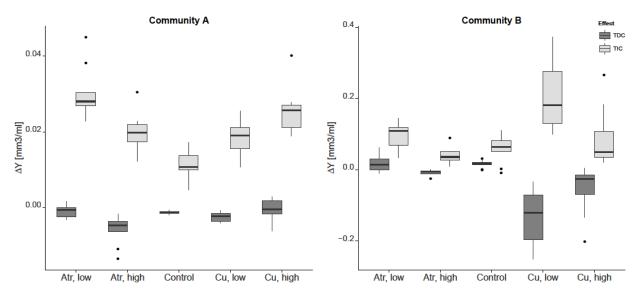
In community A, all tested levels of atrazine and copper increased the two biodiversity effects (Fig.
4.1A). This increase drove the small and negative dominance effect closer to zero and made the
large and positive complementarity effect more positive (Fig. 4.1A, Addendum III Tables S3 and
S4).

In community B, both stressors also changed complementarity and dominance (Fig. 4.1B).
However, in contrast to community A, these changes were negative at high stress levels. The small
and positive dominance effect was reduced by all stress levels, driving it closer to zero (Fig. 4.1B,
Addendum III Table S3). The large and positive complementarity effect was reduced by high levels
of both stressors, but was still well above zero (Fig. 4.1B, Addendum III Table S4).



2104 2105 **Fig. 4.1:** Dominance and complementarity effects in two benthic diatom communities exposed to low and 2106 high (200 and 500  $\mu$ g/L) concentrations of copper (Cu) and atrazine (Atr). ΔY is the deviation in the 2107 observed from the expected yield of diatom communities that could be attributed to either dominance and 2108 complementarity effects. Complementarity is indicated as the sum of two-way and one-way 2109 complementarity effects.

2111 Complementarity effects in community A were essentially composed of two-way 2112 complementarity (TIC, Fig. 4.2A), which was consequently always positive and positively affected 2113 by all stress levels (Fig. 4.2A, Addendum III Table S5). The contribution of two-way and one-way 2114 complementarity (TDC) were more comparable in community B than in community A. Complementarity effects in community B were mostly composed of TIC (control and atrazine 2115 2116 treatments), and by both TDC and TIC (copper treatments, Fig. 4.2B). Interestingly, low stress had 2117 no (atrazine) or negative (copper) effects on TDC and positive (copper) or no (atrazine) effects on 2118 TIC, while high levels of both stressors had negative effects on TDC (Fig. 4.2B, Addendum III Table 2119 S5). Thus, negative effects on TDC were partly offset by positive effects on TIC when stress was 2120 low, but not when stress was high.



**Fig. 4.2:** Two-way complementarity effects (TIC) and one-way complementarity effects (TDC) in two benthic diatom communities (A and B) exposed to low and high (200 and 500  $\mu$ g/L) concentrations of copper and atrazine.  $\Delta$ Y is the deviation in the observed from the expected yield of diatom communities that could be attributed to either TIC or TDC.

2127 Linking stress effects on two-way complementarity to EPS production. The release of EPS 2128 predicted the increase of TIC in community A under atrazine and in community B under copper 2129 (Fig. 4.3). In community A, diatoms increased their EPS production at low atrazine and copper 2130 stress. EPS secretion and TIC were positively related when diatoms were exposed to atrazine (Fig. 2131 4.3A, Addendum III Table S6), but not to copper. In community B, diatoms increased their EPS 2132 production at low copper, but not atrazine stress. EPS and TIC were not related under atrazine 2133 stress, but the amount of EPS produced by the diatoms positively affected TIC in communities 2134 exposed to copper (Fig. 4.3B, Addendum III Table S6).

## 2135 Linking stress effects on one-way complementarity to stress tolerance and monoculture

2136 **yield.** In community A, the relative performance of diatoms (the deviation of the observed relative

2137 yield from the observed frequency in community, the main component of TDC) was not related to

2138 stress tolerance or monoculture yield (Fig. 4.4 A1-A2, Addendum III Table S7). Diatoms in this

community differed in both their stress tolerance and monoculture yield (see x-axes in Fig. 4.4 A1-

- A2), but growth in community did not benefit species with either above- or below-average stress
- tolerance or monoculture yields.
- 2142 In community B, atrazine tolerance and monoculture yield also did not predict the relative
- 2143 performance of diatoms in communities exposed to atrazine (Fig. 4.4 B1-B2, Addendum III Table
- S7). Under copper stress, the relative performance of diatoms was negatively related to both their
- 2145 copper tolerance and monoculture yield (Fig. 4.4 B1-B2, Addendum III Table S7). These negative

2146 correlations were caused by *Amphora lineolata*, *Gyrosigma* sp. 1 and *Navicula digitoradiata*, which 2147 were the least copper-tolerant and lowest-yield-species within community B (Fig. 4.4 B1-B2). Despite their low monoculture yields and stress tolerance, *N. digitoradiata, Gyrosigma* sp. 1, and 2148 2149 A. lineolata however showed the highest relative performance when grown in community (Fig. 4.4 2150 B1-B2). In community B one-way complementarity under copper stress thus benefited the species 2151 which were most sensitive to the metal, but also had the lowest biomass yields, which led to a 2152 decline of total complementarity and ultimately caused a decrease of the net biodiversity effect 2153 on diatom biomass under copper stress.

2154

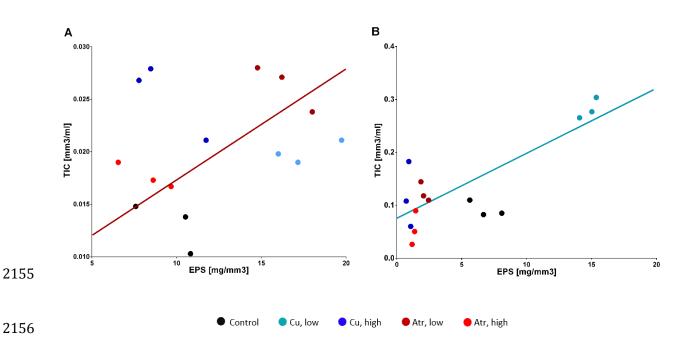
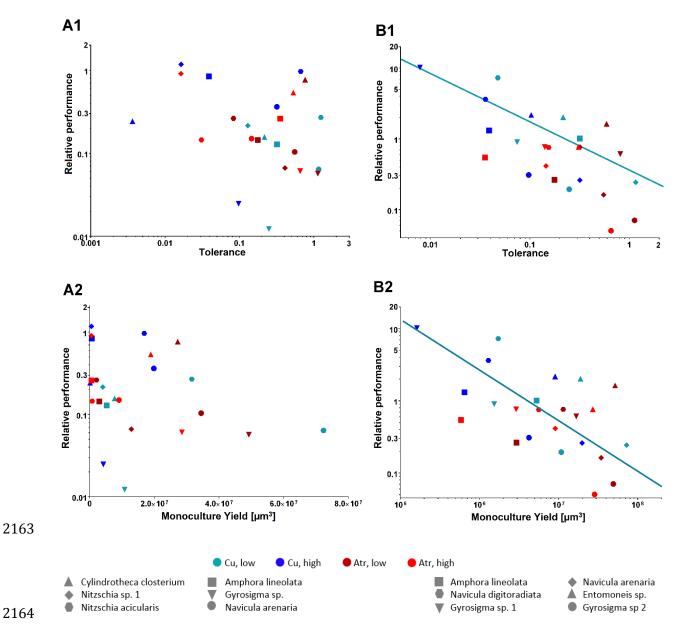


Fig. 4.3: Two-way complementarity (TIC) as a function of the release of extracellular polymeric substances
(EPS) by two benthic diatom communities (panels A and B) exposed to low and high (200 and 500 μg/L)
concentrations of copper (Cu) and atrazine (Atr). Regression lines show significant relations between EPS
and TIC in diatom communities when exposed to copper (blue line) and atrazine (red line). Regressions
were fitted over three treatments (control, 200 and 500 μg/L) for each stressor.

2162



2165 Fig. 4.4: Relative performance of diatom species in community as a function of their stress tolerance and 2166 biomass yield in monoculture. The panels A1 and B1 show the relative performance of diatom species 2167 (calculated as the deviation of observed relative yield (RY) from the relative frequency (RY/RYT) as a 2168 function of stress tolerance (calculated as the ratio of the monoculture yield at each stress level and in the 2169 control) in community A (panel A1) and B (panel B1). Panels A2 and B2 show the relative performance as a 2170 function of monoculture yield for diatoms in community A (panel A2) and B (panel B2). Stress levels are 2171 low and high (200 and 500 µg/L) atrazine (Atr) and copper (Cu). Regression lines show significant relations 2172 between the deviation of the species' relative performance in community and their stress tolerance and 2173 biomass yield in monoculture. Data are shown on a log scale, as models were fitted to log-transformed data.

#### 2175 **4.4. Discussion**

The outcome of our experiment shows a consistently positive diversity effect on diatom biomass
production which was mainly caused by complementarity rather than dominance effects. The
diatoms' production of extracellular polymers predicted, in part, increases in two-way

complementarity. These increases could however be offset if complementarity was one-way:
when low-yield species were least stress-tolerant in monoculture but benefited most from
complementarity, the diversity effect on the biomass yield of stressed diatom communities was
reduced.

2183 Diatom biomass production is crucial for the functioning of muddy intertidal systems, since 2184 benthic diatoms represent the main primary producers and thus the basis of the food web in these 2185 areas (Underwood and Kromkamp 1999, Tang and Kristensen 2007). Our findings of 2186 complementarity as the main driver of diatom biomass corresponds to previous works which also 2187 found complementarity rather than dominance to cause positive diversity effects on microalgae 2188 biomass yield (Vanelslander et al. 2009, Cardinale 2011, Stockenreiter et al. 2012, Schabhüttl et 2189 al. 2013). More importantly, these complementarity effects increased when diatoms were 2190 exposed to pesticide and metal stress, highlighting the value of producer diversity for maintaining 2191 biomass production at the basis of intertidal food webs under stressful conditions. Changes in 2192 environmental conditions can increase complementarity effects on primary producer biomass 2193 yield if facilitation, one of the mechanisms potentially leading to complementarity, becomes more 2194 important under harsh environmental conditions (Wang et al. 2013, Hodapp et al. 2016).

2195 The increase of two-way complementarity was, in part, predicted by a stimulation of diatom EPS 2196 production under stress. The release of these extracellular polymers is a common stress response 2197 in diatom biofilms: diatoms can increase their EPS production in response to nitrogen depletion, 2198 low temperatures or other unfavourable abiotic conditions (Staats et al. 2000a, Wolfstein and Stal 2199 2002). EPS production has furthermore been linked to protection from desiccation, physical 2200 disturbance and resistance to toxins (Decho 1990, Gerbersdorf et al. 2009b). Here, diatom 2201 communities increased their EPS production at low stress levels, thus confirming the stimulation 2202 of EPS as a response to low stress, which can however not be maintained under highly stressful 2203 conditions. This corroborates previous findings, which identified the release of organic exudates 2204 as a protective mechanism against chemical stress (Pistocchi et al. 1997), which can also lead to 2205 facilitation and thus increased complementarity in diatom biofilms (Vanelslander et al. 2009).

Diatoms in community B did not increase their EPS production under atrazine stress, and there was no relation between EPS production and complementarity, which did not change under atrazine. In community A however, a stimulated EPS production predicted the increase of complementarity under atrazine. Atrazine blocks the electron transport chain of Photosystem II (PSII, Dorigo and Leboulanger 2001). Some diatom species can however reduce their dependency on photosynthesis and thus their sensitivity to PSII inhibitors by mixotrophic growth, i.e. the uptake of organic substances, if such substances are present in sufficient amounts in the medium

2213 (Debenest et al. 2009, Larras et al. 2014). Amongst the diatom species in community A, 2214 *Cylindrotheca closterium* has been shown to increase its growth in response to organic exudates 2215 by other diatom species (Vanelslander et al. 2009), and C. closterium, Nitzschia sp. 1 and Nitzschia 2216 acicularis can grow in the presence of atrazine when the growth medium contains organic carbon 2217 sources (Chapter 3). When community A was exposed to atrazine, these three species showed the 2218 highest relative performance (deviation of observed relative yield from observed frequency, see 2219 y-Axis Fig. 4.4 A1). Although our findings show that the mixotrophic species in community A 2220 performed best under atrazine, one-way complementarity stayed close to zero. This was due to 2221 the – compared to the other species in community A – average monoculture yields of these best-2222 performing mixotrophic species, which resulted in neither positive nor negative covariances 2223 between the species' relative performance and monoculture yields, and thus only minor one-way 2224 complementarity effects. The increase of diatom EPS production under atrazine stress thus served 2225 as a potential facilitative mechanism, which drove complementarity effects through the presence 2226 of mixotrophic species capable of utilizing these organic exudates. Since these mixotrophic species 2227 were not characterised by any above- or below average monoculture yields, the stimulation of EPS 2228 production led to an increase of two-way rather than one-way complementarity.

2229 Both diatom communities also increased their EPS production when exposed to copper. 2230 Microalgae are usually among the first organisms to be affected by metal pollution due to their 2231 large surface to volume ratio (Miao et al. 2005, Quigg et al. 2006, Manimaran et al. 2012). The 2232 toxicity of heavy metals to microalgae primarily lies in the formation of reactive oxygen species, 2233 which can damage cell membranes, plastids and other intracellular structures (Masmoudi et al. 2234 2013). However, microalgae can increase their metal tolerance through the excretion of 2235 extracellular polymers (Hall et al. 1979, Fisher and Frood 1980, Pistocchi et al. 1997). The 2236 excretion of EPS allows microalgae to chelate metal ions in the medium, thus reducing metal 2237 accumulation within the algal cells (Pistocchi et al. 1997). Compared to other microalgal species, 2238 diatoms are characterised by a high EPS production (De Brouwer et al. 2000). This capacity to 2239 excrete EPS under metal stress can enable diatoms to grow at copper concentrations which are 2240 up to one magnitude higher than those limiting the growth of other microalgae classes (Pistocchi 2241 et al. 1997).

In community A, most species performed better when exposed to copper in community rather than in monoculture, but this complementarity was evenly distributed amongst species, and not predicted by their EPS production. The positive diversity effect on the biomass yield of this diatom community under copper was thus likely caused by other mechanisms than the excretion of extracellular polymers. Such mechanisms can consist of an intracellular accumulation of metal ions in vacuoles or metal immobilization in the cell wall, selective copper exclusion from algal cells, and an increased production of amino acids protecting cell membranes from copper damage(Fernandes and Henriques 1991, Lage et al. 1994, Wu et al. 1995).

2250 The diatoms' EPS production predicted complementarity in community B under copper stress. 2251 However, this complementarity was not equally distributed amongst species, and essentially 2252 favoured A. lineolata, Gyrosigma sp. 1, N. digitoradiata. These species had the lowest biomass 2253 yields and were least copper-tolerant in monoculture, but showed the highest relative 2254 performance when exposed to the metal in community. These high relative performances did not 2255 come at the expense of other species (neutral dominance effect), but were due to a one-way 2256 complementarity which only benefited sensitive species with low monoculture yields, causing the 2257 increases in two-way complementarity to be mirrored and offset by a negative one-way 2258 complementarity. Diversity thus increased the biomass yield of diatom species under copper 2259 stress, but since sensitive and low-yield species benefited most from being grown in community, 2260 the net diversity effect on diatom biomass yield was not altered or even reduced at low and high 2261 copper stress respectively.

2262 Complementarity effects tend to increase over time (Fargione et al. 2007). Also, complementarity 2263 is strongest along gradients of broad environmental niche space, which allow for resource 2264 partitioning between species under natural conditions (Spaèková and Lepš 2001, Hodapp et al. 2265 2016). Our experiments were performed over a relatively short time (one diatom growth cycle) 2266 in an artificial experimental system which limited environmental fluctuations and spatial 2267 heterogeneity, which can lead to an underestimation of complementarity effects. Also, to confirm 2268 EPS excretion as a facilitative mechanism under stress, it is recommended to quantify the extent 2269 to which extent the production of EPS reduced metal exposure of diatom cells, as well as the 2270 uptake of these organic exudates by mixotrophic species. The former could be achieved by 2271 measuring the intracellular copper concentrations in the diatoms (e.g. see Pistocchi *et al.* 1997), 2272 the latter by a colorimetric quantification of the metabolization of organic substances by diatoms 2273 (e.g. see Tuchman et al. 2006). Moreover, EPS production was a 'sleeping' facilitative mechanism, 2274 which was stimulated when diatom communities were exposed to chemical stress, and only 2275 predicted complementarity in diatom communities containing either stress-sensitive species, or 2276 species capable of heterotrophic growth.

We infer four conclusions from our results: first, a positive biodiversity effect causes diatom communities to produce more biomass than expected from their monoculture. Second, this positive biodiversity effect is largely underpinned by complementarity effects. Third, complementarity in stressed diatom communities can be driven by the release of extracellular polymers as a facilitative mechanism, leading to a better performance of species characterised by

- 2282 a mixotrophic growth mode or a low stress tolerance in monoculture. Fourth, a negative one-way
- 2283 complementarity can cause the net diversity effect to decrease under stress, if low-yield species
- 2284 perform better when grown with different species.

# Chapter 5: Selective and context-dependent effects drive chemical stressor impacts across trophic levels at the basis of marine food webs

2288

#### 2289 Abstract

Human activities increasingly impact the functioning of marine food webs, but anthropogenic
stressors are seldom included into ecological study designs and diet quality, as distinct from just
diet quantity, has rarely been highlighted in food web studies in a stress context.

2293 We first measured the effects of metal and pesticide stress (copper and atrazine) on the 2294 contribution of a benthic intertidal diatom community to three processes that are key to the 2295 functioning of intertidal systems: biomass production (diet quantity), energy content (diet 2296 quality) and extracellular polymer production (sediment stabilization). We then examined if 2297 stressors affected diatom functioning by selectively targeting the species contributing most to 2298 functioning (selective stress effects) or by changing the species' functional contribution (context-2299 dependent effects). Finally, we tested the response of the diatoms' main grazers (harpacticoid 2300 copepods) to changes in diet quality.

Diatom diet quantity was reduced by metal stress but not by low pesticide levels due to the
presence of an atrazine-tolerant, mixotrophic species. Context-dependent effects of both stressors
increased the diatoms' contribution to sediment stabilization by stimulating the release of
extracellular polymers.

Selective effects of the pesticide reduced diatom diet quality by 60% and 75% at low and high pesticide levels respectively, by shifting diatom community structure from dominance by lipidrich species towards dominance by atrazine-tolerant, but lipid-poor species. Context-dependent effects did not affect individual diatom lipid content at low levels of both stressors, but caused diatoms to lose 40% of their lipids at high copper stress.

Copepod energy content was related to the quality of their diatom diet, with copepods losing half of their energy content (in terms of lipids) when feeding on diatoms grown under low and high pesticide and high metal stress. Selective atrazine effects on diatom community structure affected the energy flow to their grazers at stress levels where no context-dependent effects of both stressors on diatom diet quality were detected.

*Synthesis.* Chemical stress differentially affected the contribution of marine primary producers to
diet quantity, sediment stabilization and diet quality. Selective community shifts were a more
potent threat for diet quality than context-dependent stress effects on primary producers. Diet

quality predicted the energy flow from marine producers to their consumers and should beconsidered when measuring food web functioning under anthropogenic change.

2320

### 2321 **5.1. Introduction**

The impact of human activities on biological communities and their contribution to ecosystem functioning has become a central topic in ecological research (Halpern et al. 2008, Cardinale et al. 2012, Gamfeldt et al. 2015). Although conservation research is framed within the context of anthropogenic change, exposure to anthropogenic stressors is rarely included into ecological study designs (McMahon et al. 2012, Steudel et al. 2012).

2327 Stress can affect ecosystem functioning by causing biodiversity loss in terms of species richness, 2328 as well as through changes in community structure, without necessarily causing species to go 2329 extinct (Hillebrand et al. 2008, Wittebolle et al. 2009, Mensens et al. 2015). Selective stress effects 2330 on community structure (hereafter 'selective stress effects', Wittebolle et al. 2009) can reduce 2331 ecosystem functioning if stressed communities are dominated by tolerant species with a low 2332 functional contribution (Larsen et al. 2005, Mensens et al. 2015). If the functionally most 2333 important species are also most stress-tolerant, loss of functioning under stress will be limited 2334 (Radchuk et al. 2016). Moreover, functioning in stressed communities can be driven by 'context-2335 dependent effects', i.e. changes in the species' functional contribution (Fox 2006, Fox and Harpole 2336 2008, Tylianakis et al. 2008, Hiddink et al. 2009). Context-dependent effects can arise from direct 2337 effects of environmental drivers on the species' functional contribution (Fox and Harpole 2008, 2338 e.g. physiological stress, Schimel et al. 2007), as well as from differential species interactions 2339 under stress (e.g. changes in complementarity, Fox 2006, Fox and Harpole 2008).

2340 The majority of experiments designed to address ecosystem functioning under anthropogenic 2341 change have focused on single trophic levels, usually primary producers (Raffaelli 2006, Cardinale 2342 et al. 2011). Stressors that alter functioning at the producer level can however have concomitant 2343 impacts on their consumers (Rohr and Crumrine 2005, McMahon et al. 2012). The trophic impacts 2344 of anthropogenic stressors are commonly examined with regard to diet quantity, which has been 2345 linked to the abundance and biomass of consumers (e.g. Kasai and Hanazato 1995, Wendt-Rasch et al. 2004, Rohr and Crumrine 2005). Far less attention has been devoted to diet quality, which 2346 2347 considers a diet's biochemical composition and is commonly expressed as lipid content (Jodice et 2348 al. 2006, Guo et al. 2016). Nevertheless, the growth, reproduction and energy profile of consumers 2349 is strongly linked to the quality of their diet (Giles et al. 2002, Österblom et al. 2008, Taipale et al. 2350 2013). The biochemical composition of primary producers is increasingly affected by human 2351 disturbance (Vitousek et al. 1997, Wang and Frei 2011, Guo et al. 2016, Sanpera-Calbet et al.

2352 2016), which has made the integration of food biochemistry with traditional studies of diet
2353 quantity a key challenge for estimating food web functioning under stress (Arts and Wainmann
2354 1999, Guo et al. 2016).

2355 Here, we first measure the effects of metal and pesticide stress (copper and atrazine) on the 2356 structure of a benthic diatom community and its contribution to three ecosystem processes: 2357 biomass production (diet quantity), energy content (diet quality) and extracellular polymer 2358 production (sediment stabilization). Benthic diatoms are the main primary producers in many 2359 soft-sediment intertidal habitats, enhancing sediment stabilization in these systems through the production of extracellular polymeric substances (EPS, Underwood and Kromkamp 1999, Decho 2360 2361 2000). Diatom diet quality in terms of essential fatty acid (EFA) content plays a crucial role for 2362 trophic energy transfer (Arts et al. 2001, Taipale et al. 2013). EFAs cannot be synthesized by 2363 animals but are key determinants of the growth and energy content of aquatic consumers (Brett 2364 and Müller-Navarra 1997, von Elert 2002, Arendt et al. 2005, Litzow et al. 2006). Harpacticoid 2365 copepods are among the main consumers (grazers) of benthic diatoms and incorporate large 2366 amounts of EFAs from their algal diet, making harpacticoids pivotal for the energy transfer from 2367 primary producers to higher trophic levels (Alheit and Scheibel 1982, De Troch et al. 1998, Nanton 2368 and Castell 1998, Buffan-Dubau and Carman 2000, Andersen et al. 2005).

2369 Next, we test if stressor-induced changes in diatom functioning (EFA and EPS production) were 2370 caused by selective or context-dependent stress effects. Context-dependent effects are measured 2371 by comparing functioning in experimental diatom communities with that in synthetic 2372 communities. Synthetic communities are computed from the species' EFA and EPS production in 2373 unstressed monocultures and reflect the same community structure observed at each stress level, 2374 thus allowing to exclude any confounding effects of community structure on diatom functioning. 2375 Selective stress effects are quantified by contrasting synthetic communities which reflect the 2376 community structure under stress and control conditions, thus excluding any confounding 2377 context-dependent effects on diatom functioning.

Last, we test if potential changes in the survival and energy content of the harpacticoid copepod *Microarthridion littorale*, being the dominant grazer at the intertidal study site, are related to atrazine and copper effects on the quality of its diatom diet. For instance, whether potential changes in grazer energy content are linked to context-dependent stress effects on diatom energy content, or to changes in community structure leading to dominance of diatoms with a low or high contribution to energy flow in intertidal systems.

#### 2384 **5.2. Methods**

2385 Experimental organisms & culture conditions. The harpacticoid copepod Microarthridion 2386 littorale (family Tachidiidae) was collected from intertidal mud at the Paulina intertidal flat (SW 2387 Netherlands, 51° 21'N, 3°43'E), where it represented the dominant grazer (~ 90% of all 2388 harpacticoid individuals). M. littorale specimens were extracted alive from the sediment using a 2389 mixed technique of sediment decantation and extraction via white light attraction. Adult 2390 specimens were randomly collected with a glass pasteur pipette using a Wild M5 binocular. 2391 Copepods were washed 3 times over a 38 µm sieve and placed in glass jars with filtered and 2392 autoclaved natural seawater overnight in order to empty their intestines prior to the start of the 2393 experiment.

2394 The diatom community was composed of six species representing the most abundant genera 2395 observed at the sampling site (Addendum IV Table S1). For this experiment, all species were 2396 obtained from the culture collection of the Protistology and Aquatic Ecology Research Group 2397 (UGent) (http://bccm.belspo.be). Prior to the set-up of the experiments, the diatoms were grown 2398 in tissue bottles (Greiner BioOne, CELLSTAR® TC, 175 cm<sup>2</sup> growth surface) during 10 days in a 2399 climate room at 15±1 °C, a light/dark cycle of 12h / 12h and an illumination of 90 µmol photons 2400 m<sup>-2</sup>s<sup>-1</sup>, in culture medium consisting of filtered and autoclaved natural seawater (salinity 32±1) 2401 enriched with f/2 nutrients (Guillard 1975).

2402 **Diatom experiments.** The experimental diatom communities were exposed in five treatments to 2403 0, 200 (hereafter 'low') and 500 (hereafter 'high')  $\mu$ g/l atrazine and copper respectively, 2404 representing levels of atrazine and copper stress resulting on average in a 50% and 90% 2405 reduction in growth of the diatom species (Chapter 3). Atrazine treatments were prepared from a 2406 stock solution obtained by dissolving 50 mg technical atrazine (2-chloro-4-ethylamino-6-2407 isopropylamino-s-triazine, 99.8% pure, Sigma-Aldrich Chemie Gmbh, Munich, Germany) in 10 ml 2408 acetone as a carrier to increase the solubility of atrazine, with a maximum final volume of 0.01% 2409 acetone in the treatments. An acetone control treatment of 0.01% acetone was included and 2410 compared to an acetone-free control to test for carrier effects. All atrazine treatments were 2411 compared to the acetone control. Copper (as a Cu[II]Cl<sub>2</sub> solution, analytical grade; VWR 2412 International) was spiked directly into the culture medium before exposure of the diatoms. F/2 2413 culture medium for the copper experiments was prepared without EDTA, to avoid complexation 2414 of free copper ions (Pistocchi et al. 1997). The obtained atrazine and copper concentrations that 2415 were finally applied in the experiment are listed in Addendum IV Table S2. Additionally, the six 2416 diatom species were assembled in monoculture under control conditions, to quantify each 2417 species' biomass, EFA and EPS production in the absence of the stressors (Addendum IV Table 2418 S1). All treatments were run in tissue culture flasks (Greiner BioOne, CELLSTAR® TC, 175 cm<sup>2</sup>

2419 growth surface), with nine replicates per treatment (three replicates in the monoculture 2420 treatments). Each microcosm was inoculated with a total cell density of approximately 5000 2421 diatom cells/ml (500000 cells in 100 ml medium per culture flask, between 800-850 cells/ml per 2422 species in the diatom communities) from exponentially growing cultures, and incubated in a 2423 climate room at 15±1 °C, under a light / dark cycle of 12h / 12h at 90 µmol photons m<sup>-2</sup>s<sup>-1</sup>. Culture medium was refreshed after eight and 15 days, diatom biofilms as food for the copepod 2424 2425 experiment as well as for the EPS and EFA analyses (see further) were harvested after 15 days 2426 (late exponential growth phase) and experiments were terminated after 25 days.

2427 Diatom biofilms in three replicates of 100 ml cell suspension per treatment were harvested as 2428 food for the corresponding treatments in the copepod experiment, and purified from copper and 2429 atrazine. In the purification process, the diatom cells were resuspended and placed in 50 ml 2430 centrifuge tubes (2 tubes per replicate). All samples were then centrifuged at a speed of 50g for 2431 10 minutes. Supernatant fluid was removed and replaced with f/2 culture medium (25 ml total 2432 volume) and centrifuged again for 10 minutes. After the second centrifuge process, supernatant 2433 fluid was removed again, and 5 ml of concentrated diatom pellet per replicate were transferred to 2434 nine Eppendorf microtubes (0.56 ml pellet per microtube), freeze-dried and preserved at -80 °C. 2435 The individual microtubes contained the food aliquot for each day of the respective treatments in 2436 the copepod experiment (see further).

2437 **Copepod experiment.** We tested the effect of diatom diet quality by offering *M. littorale* diatom 2438 diets of equal biomass under unstressed conditions. The copepod experiment consisted of five 2439 treatments, each with three replicates of 100 *M. littorale* copepods (a natural mix of adult males 2440 and (gravid) females), that were fed for 10 d an equal biomass of diatoms grown under unstressed 2441 conditions and low and high atrazine and copper stress, respectively. The experiment was 2442 conducted in glass jars containing 100 ml of filtered and autoclaved seawater in a climate room at 2443 15±1 °C with a 12:12 h light:dark cycle and 40 to 50 µmol photons m<sup>-2</sup> s<sup>-1</sup>. To ensure a constant 2444 food supply, each treatment was inoculated with a concentrated diatom pellet of 0.23 mm<sup>3</sup> 2445 biovolume every day, and unconsumed diatoms were removed from the bottom of the jars. Over 2446 the whole duration of the experiment, a total diatom biovolume of 2.05 mm<sup>3</sup> was applied per 2447 experimental unit, corresponding to 3 to 5 x 10<sup>6</sup> diatom cells per treatment. At the end of each day, 2448 there was no food depletion in any of the treatments. Based on our previous experiments (De 2449 Troch et al. 2005, 2007), the provided quantity of diatoms can be considered as above the feeding 2450 saturation level. Copepod mortality was determined at the end of the experiment, and surviving 2451 (85-100%) individuals from each experimental unit were washed in natural seawater to remove 2452 food particles, left 12h to empty their gut, and stored at – 80 °C for further fatty acid analysis.

2453 Biomass, EPS and EFA analyses. Diatom biomass was quantified as biovolume after 0, 2, 5, 10, 2454 15, 20 and 25 days of incubation. Biovolume was calculated from cell densities, linear dimensions 2455 (measured digitally using ImageJ, Schneider et al. 2012) and formulas representing the closest 2456 approximation of geometric shape for each genus (Hillebrand et al. 1999, Addendum IV Table S1). 2457 Cell densities (in cells/ml) were determined by digitally counting the cells (ImageJ cell counting 2458 software) in photographs obtained by magnifying and photographing (x100) an area of 0.66 mm<sup>2</sup> 2459 per microcosm, using an inverted Axiovert 135 Zeiss microscope (Carl Zeiss, Jena, Germany) and 2460 a connected digital camera (Canon PowerShot G11). All further analyses use the biomass on day 2461 15.

Extracellular polymeric substances (EPS) secreted by the diatom communities were measured in three replicates of 10 ml diatom suspension which were collected after 15 days of incubation. EPS concentrations were analysed by spectrophotometry according to a modified version of the phenol/H<sub>2</sub>SO<sub>4</sub> assay (Dubois et al. 1956) as described in Mensens *et al.* (2015).

Essential fatty acids (EFAs) as marker of energy content were quantified as the content of eicosapentaenoic acid ( $20:5\omega3$ , EPA) and docosahexaenoic acid ( $22:6\omega3$ , DHA). Three replicates of 8 ml suspended diatom culture per treatment of the diatom communities and monocultures were collected after 15 days. The samples were centrifuged for 10 minutes at 10 °C and a speed of 50*g*. After undergoing the same purification process as the diatom food samples for the grazer experiment, pellets were resuspended and 1.5 ml of concentrated diatom suspension per replicate were placed in a glass vial and stored at -80 °C for fatty acid analysis.

2473 Copepod and diatom EFA content was measured through hydrolysis of total lipid extracts and 2474 methylation to FA methyl esters (FAME), followed by the analysis of the obtained FAME using a 2475 gaschromatograph (HP 6890N) coupled to a mass spectrometer (HP 5973) according to the 2476 protocol described in De Troch et al. (2012) for copepods and Mensens et al. (2015) for diatoms. 2477 The quantification function of each individual FAME was obtained by linear regression of the 2478 chromatographic peak areas and corresponding known concentrations of the standards (ranging 2479 from 5 to 250 ng/ml). All EPS and EFA concentrations were standardized to diatom biomass or 2480 the number of copepod individuals for diatoms and copepods, respectively.

Data analysis. An analysis of the diatom community structure among the treatments (control, low and high atrazine and copper) was conducted with a non-metric multidimensional scaling method based on Bray-Curtis similarity. A one-way analysis of similarity (ANOSIM) was used to test for significant biomass differences between the treatments. Subsequently, percentages of similarity (SIMPER) were calculated to determine the main species contributing to any differences in community structure. Differences in diatom biomass, EFA and EPS production among the treatments were tested with a generalized least squares model, with biomass, EFA and EPS as response variables and treatment type as categorical predictor (Equation 1)

2490  $Y \sim \beta_T \cdot T$ 

[1]

2491 where Y is the response variable (biomass, EFA or EPS), T is the treatment type (control, low and 2492 high atrazine and copper), and the beta coefficient  $\beta_T$  is the slope measuring the effect of the 2493 treatment type on biomass, EFA or EPS production. Biomass, EFA and EPS production among 2494 treatments were compared with pairwise Tukey's tests correcting p-values for multiple 2495 comparisons by the single-step method.

2496 Next, we tested if potential changes in diatom EFA and EPS production were due to selective or 2497 context-dependent effects of atrazine and copper. Selective and context-dependent effects were 2498 quantified by comparing the EFA and EPS production in experimental and synthetic diatom 2499 communities. The synthetic communities have the same community structure as observed in the 2500 corresponding treatments of the experimental communities, but are computed from the species' 2501 EFA and EPS production in unstressed monocultures (Equation 2). The synthetic communities 2502 thus reflect the EFA and EPS production expected at the same community structure as induced by 2503 copper and atrazine, however without any stress exposure or species interactions.

$$Y_{SYN,j} = \frac{\sum_{i=1}^{N} M_{i,j=0} \times B_{i,j}}{B_{T,j}}$$
[2]

2504  $Y_{SYN,j}$  is the EFA or EPS production in synthetic communities of the same structure as the 2505 experimental communities at the atrazine or copper concentration *j*.  $M_{i,j=0}$  is the mean EFA or EPS 2506 production per unit biomass of species *i* in monoculture under unstressed conditions (*j*=0).  $B_{i,j}$  is 2507 the biomass of species *i* observed in the experimental community at the stress level *j*.  $B_{T,j}$  is the 2508 total biomass observed in the experimental community at the stress level *j*.

2509 Selective and context-dependent effects of both stressors were analysed with a generalized least 2510 squares model (Equation 3) and pairwise comparisons of the EFA and EPS production in 2511 experimental and synthetic diatom communities.

$$2512 \quad Y \sim \beta_{\rm T} \cdot {\rm T}$$
 [3]

where Y is the diatom EFA or EPS production per unit biomass, T is the treatment type (control, low and high atrazine and copper in the experimental and synthetic diatom communities), and the beta coefficient  $\beta_T$  is the slope measuring the effect of the treatment type on EFA or EPS production. Pairwise comparisons were performed with a Tukey's test correcting p-values formultiple comparisons by the single-step method.

2518 Context-dependent effects occur when synthetic and experimental communities within the same 2519 treatment differ in their EFA or EPS production. Since both community types have the same 2520 structure, any differences in EFA or EPS between the two community types result from direct 2521 stress effects or species interactions in the experimental community. Consequently, any 2522 differences between experimental and synthetic communities of the same treatment point at 2523 context-dependent effects.

2524 Selective stress effects occur when synthetic communities reflecting the control community 2525 structure differ in their EFA or EPS production from synthetic communities reflecting the 2526 community structure under stress. Since synthetic communities are computed from unstressed 2527 monocultures, any differences in EFA or EPS between synthetic communities are related to 2528 differences in community structure rather than direct stress effects or species interactions. 2529 Consequently, any differences among synthetic communities are linked to selective rather than 2530 context-dependent effects. Addendum IV Fig. S1 provides a scheme visualizing the quantification 2531 of context-dependent and selective stress effects.

2532 The response of copepod energy content to stressor-induced alterations in diet quality was 2533 analysed with generalized least squares models, with copepod energy content as response 2534 variable and diatom diet quality in terms of energy content and community structure as predictors 2535 (Equation 4). Diatom energy content was quantified as EFA production, diatom community 2536 structure as the Bray-Curtis percent dissimilarity from the average community structure in 2537 controls (Equation 5). Models were fitted separately for copepods feeding on atrazine- and 2538 copper-exposed diatoms respectively, to test if the effects of either stressor on copepod energy 2539 content can be predicted from changes in diatom energy content or community structure.

$$2540 \qquad E_{\rm C} \sim a + b \cdot E_{\rm D} + c \cdot C_{\rm D} \tag{4}$$

2541  $E_c$  is the copepod energy content (EFA content per copepod individual),  $E_D$  is diatom energy 2542 content (EFA production per unit biomass),  $C_D$  is diatom community structure, a is the intercept, 2543 b and c represent the slopes, i.e. the relation of copepod energy content to diatom energy content 2544 and diatom community structure. If  $E_D$  and  $C_D$  were correlated (correlation factor > 0.5), models 2545 were fitted separately for both predictors.

2546 
$$C_D = 100 \cdot \{1 - \Sigma | B_{i,j} - \mu B_{i,j=0} | / \Sigma (B_{i,j} + \mu B_{i,j=0})\}$$
 [5]

2547  $B_{i,j}$  is the biomass of species *i* at the atrazine or copper concentration *j* and  $\mu B_{i,j=0}$  is the mean 2548 biomass of species *i* in the control (*j*=0).

2549 For all least squares model fits, normality and homogeneity of model residuals were inspected by 2550 evaluation of quantile-quantile plots and Shapiro-Wilk's test, and by Levene's test and plotting 2551 residuals versus explanatory variables respectively. Untransformed data did not violate normality 2552 (Shapiro-Wilk's test,  $\alpha > 0.1$ ). If indications of deviations from normality were detected ( $0.1 < \alpha < 0.1$ ). 2553 0.15), an optimal Box-Cox transformation (Box and Cox 1964, Venables and Ripley 2002) was 2554 applied to maximize normality of model residuals. If homogeneity was violated, the model was 2555 refitted using an exponential variance structure allowing residuals to change with the continuous 2556 predictor X (weights = varExp(form  $\sim 1 | X$ ) or allowing different variances according to the 2557 categorical predictor P (weights = varIdent(form  $\sim 1 \mid P$ ). By means of likelihood ratio testing, the 2558 significance of these structures was tested ( $\alpha = 0.05$ ).

2559 Multivariate, ANOSIM and SIMPER analyses of diatom community structure were performed 2560 using Primer 6 software (Clarke and Gorley 2006). All other calculations were done in R 3.0.1. using RStudio (R Development Core Team 2016). The package nlme (Pinheiro et al. 2015) was 2561 2562 used for the fitting of generalized least squares models and optional variance structures. Optimal 2563 Box-Cox transformations were performed using MASS (Venables and Ripley 2002). Pairwise 2564 Tukey's tests on the fitted models were performed with the package multcomp (Hothorn et al. 2565 2008), using the general linear hypothesis test (glht) function, correcting p-values for multiple 2566 comparisons by the single-step method (default procedure in multcomp).

2567

#### 2568 **5.3. Results**

2569 Diatom community structure. The structure of diatom communities under atrazine differed 2570 from the structure of communities grown under control conditions and copper exposure (see non-2571 metric multidimensional scaling in Addendum IV Fig. S2). The ANOSIM confirmed the significance 2572 of treatment type for community structure (global R = 0.833, p=0.001). The community structure 2573 at both copper levels resembled the community structure observed under control conditions 2574 (14% and 18% dissimilarity respectively), with N. acicularis and N. arenaria contributing the most 2575 biomass in both types of communities (Fig. 5.1, Addendum IV Table S3). In contrast, both atrazine 2576 levels induced a change in community structure (70% and 76% dissimilarity from the control) 2577 due to an increase in biomass of C. closterium, which compared to the control showed a 6- and 12-2578 fold increase in biomass in the high and low atrazine treatments, respectively. This resulted in a 2579 dominance by *C. closterium* in the atrazine-exposed communities, as it contributed more than 70% 2580 of the total biomass at both atrazine levels (Fig. 5.1, Addendum IV Table S3). Within the control, 2581 copper and atrazine treatments, the community structure of diatom communities showed little

variance (within-treatment similarities between 84% and 92%, Addendum IV Table S4).

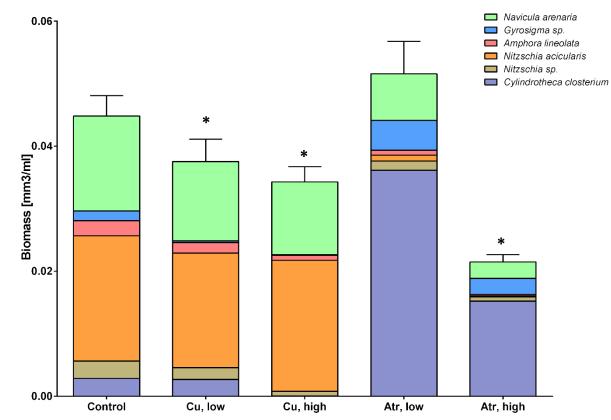


Fig. 5.1: Total biomass production per treatment and biomass of the component species for diatom communities grown in control, low (200 μg/L) and high (500 μg/L) atrazine ('Atr') and copper ('Cu') treatments. Significant differences in biomass production from the control are indicated with asterisks (\*).
2587

2588 Diatom biomass, EPS and EFA production. Diatom biomass, EPS and EFA production changed 2589 depending on treatment type (all p<0.0001, Fig. 5.1, Fig. 5.2). The post-hoc analyses showed that 2590 diatom biomass was reduced at both low and high copper as well as at high but not at low atrazine 2591 concentrations (Fig. 5.1, Table 5.1). Diatom EPS production increased at low atrazine and copper 2592 stress, with diatom communities producing twice as much EPS than under control conditions, but 2593 was not affected at high levels of both stressors (Fig. 5.2A grey bars, Table 5.1). Diatom EFA production was reduced at both levels of atrazine, but only at high copper stress (Fig. 5.2B grey 2594 2595 bars, Table 5.1). High copper stress reduced the EFA production of diatom communities by 40%, 2596 low and high atrazine stress by 60% and 75%, respectively (Fig. 5.2B grey bars).

2597

Process	Treatment	Treatment	Est	lwr	upr	Z	Р
		Atr, low	-6.756	-9.18	-4.33	2.78	0.059
	Control	Atr, high	23.327	20.90	25.75	9.61	< 0.001
Biomass		Cu, low	7.273	4.85	9.70	-3.00	0.036
DIOIIIdSS		Cu, high	10.533	8.11	12.96	4.34	< 0.001
	Atr, low	Atr, high	30.083	27.66	32.51	12.39	< 0.001
	Cu, low	Cu, high	3.260	0.83	5.69	1.34	0.666
	Control	Atr, low	-0.251	-0.44	-0.06	-4.03	< 0.001
		Atr, high	0.066	-0.13	0.26	1.06	0.984
EPS		Cu, low	-0.290	-0.48	-0.10	-4.66	< 0.001
EL2		Cu, high	0.016	-0.18	0.21	0.25	1.000
	Atr, low	Atr, high	0.317	0.13	0.51	5.09	< 0.001
	Cu, low	Cu, high	0.305	0.11	0.50	4.91	< 0.001
		Atr, low	2.143	1.30	2.99	8.00	< 0.001
	Control -	Atr, high	3.191	2.34	4.04	12.08	< 0.001
EFA		Cu, low	-0.046	-0.89	0.80	-0.17	1.000
EFA		Cu, high	1.372	0.52	2.22	5.12	< 0.001
_	Atr, low	Atr, high	0.317	0.13	0.51	5.72	<0.001

0.305

0.50

0.11

4.91

< 0.001

2599

2600 Table 5.1: Pairwise comparisons of biomass, EFA and EPS production in diatom communities as estimated 2601 by generalised least squares model fits. 'Process' indicates to which response variable models were fitted 2602 (biomass, EFA or EPS production). 'Treatment' (low (200 µg/L) and high (500 µg/L) atrazine ('Atr') and 2603 copper ('Cu')) indicates which treatments are compared. 'Est' indicates the difference in biomass (in 10<sup>3</sup> 2604 mm<sup>3</sup>/ml), EFA (in µg/mm<sup>3</sup>) and EPS (in mg/mm<sup>3</sup>) production between the compared treatments as 2605 estimated by generalized least squares models fitted to untransformed biomass and Box-Cox transformed EFA and EPS data, 'lwr' and 'upr' indicate the lower and upper confidence intervals of the estimated 2606 2607 difference. 'Z' and 'P' denote the z- and p-values corrected for multiple comparisons by the single-step 2608 method, bold values are statistically significant. 2609

Cu, high

Cu, low

2610 Selective and context-dependent stress effects on diatom EPS and EFA production. The EPS 2611 and EFA production in the experimental and synthetic diatom communities changed depending 2612 on treatment type (both p<0.0001). Pairwise comparisons of the EPS production in the 2613 experimental and synthetic communities showed no difference among the two community types 2614 in the control and the high copper and atrazine treatments (Fig. 5.2A, Table 5.2). At low levels of 2615 both stressors however, the EPS production in experimental communities was double of that 2616 expected in unstressed synthetic communities of the same structure (Fig. 5.2A, Table 5.2). Within 2617 the synthetic communities there was no difference in EPS among any of the treatments (Fig. 5.2A, 2618 Table 5.2).

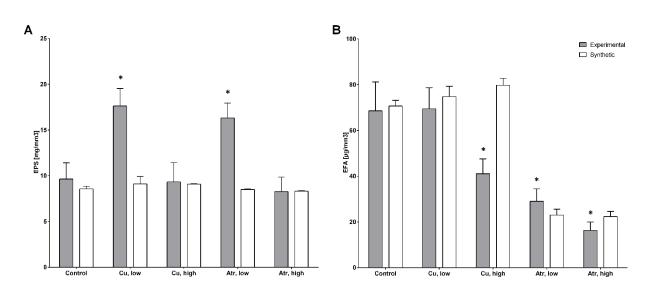
Pairwise comparisons of the EFA production in experimental and synthetic communities did not
show differences among the two community types in the control and low copper treatments (Fig.
5.2B, Table 5.2). At high copper stress, the EFA production in the experimental community was

- lower than in the corresponding synthetic community (Fig. 5.2B, Table 5.2). The EFA loss induced
- 2623 by high copper stress in experimental communities was thus not reflected in unstressed synthetic

2624 communities of the same structure, whose EFA production did not differ from the control (Fig.2625 5.2B, Table 5.2).

In the atrazine treatments, both the experimental and synthetic communities had a lower EFA production than the control (Fig. 5.2B, Table 5.2). Within each atrazine treatment, the EFA production of experimental and synthetic communities did not differ (Fig. 5.2B, Table 5.2). The EFA loss induced by atrazine in the experimental communities was thus reflected by the synthetic communities, which mimicked the community structure under atrazine without actual exposure to the herbicide (Fig. 5.2B, Table 5.2).

2632



2633

2634Fig. 5.2: Atrazine and copper effects on diatom sediment stabilization (EPS production, panel A) and energy2635content (EFA production, panel B). Experimental communities were grown in control, low (200  $\mu$ g/L) and2636high (500  $\mu$ g/L) atrazine ('Atr') and copper ('Cu') treatments. Synthetic communities were computed from2637unstressed monocultures in the same community structure observed at each stress level. Significant2638differences in the EFA or EPS production of experimental communities from the control are indicated with2639asterisks (\*).

2640

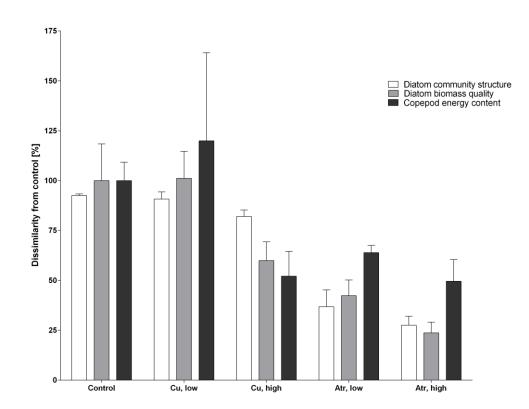
Process	Effect	Com	Treatment	Com	Treatment	Est	lwr	upr	Z	Р
			Control	Syn -	Atr, low	0.002	-0.05	0.05	0.15	1.000
	SE	Syn			Atr, high	0.012	-0.04	0.05	0.78	0.998
					Cu, low	-0.026	-0.07	0.02	-1.68	0.765
					Cu, high	-0.026	-0.07	0.02	-1.65	0.779
EPS	CD	Exp	Control	Syn	Control	0.047	-0.09	0.18	-1.05	0.984
			Atr, low		Atr, low	0.301	-0.16	0.44	6.64	< 0.001
			Atr, high		Atr, high	-0.006	-0.15	0.13	-0.13	1.000
			Cu, low		Cu, low	0.311	0.17	0.45	6.86	< 0.001
			Cu, high	-	Cu, high	0.006	-0.13	0.15	0.14	1.000
	SE	Syn	Control	Syn -	Atr, low	2.695	1.85	3.54	10.06	< 0.001
					Atr, high	2.746	1.90	3.60	10.25	< 0.001
					Cu, low	-0.173	-1.02	0.67	-0.65	0.999
					Cu, high	-0.381	-1.23	0.47	-1.42	0.921
EFA	CD	Exp	Control		Control	-0.110	-0.96	0.74	0.41	1.000
			Atr, low	nigh Syn ow	Atr, low	0.442	-0.41	1.29	1.65	0.823
			Atr, high		Atr, high	-0.554	-1.40	0.29	-2.07	0.550
			Cu, low		Cu, low	-0.237	-1.08	0.61	0.89	0.997
			Cu, high		Cu, high	-1.863	-2.71	-1.02	-6.95	< 0.001

2643 Table 5.2: Pairwise comparisons of EFA and EPS production in treatments of experimental and synthetic 2644 diatom communities as estimated by generalised least squares model fits. 'Process' indicates to which 2645 response variable models were fitted (EFA or EPS production). 'Effect' indicates which type of stress is 2646 analyzed: Selective stress effects ('SE': comparison of synthetic communities reflecting the community 2647 structure under control and stress conditions). Context-dependent stress effects ('CD': comparison of 2648 stressed and synthetic communities of the same community structure). 'Com' ('Exp': Experimental and 2649 'Syn': Synthetic) and 'Treatment' (low (200  $\mu$ g/L) and high (500  $\mu$ g/L) atrazine ('Atr') and copper ('Cu')) 2650 indicate which communities and treatments are compared. 'Est' indicates the difference in EFA and EPS production between the compared treatments as estimated by generalized least squares models fitted to 2651 2652 Box-Cox transformed EFA and EPS data, 'lwr' and 'upr' indicate the lower and upper confidence intervals of 2653 the estimated difference. 'Z' and 'P' denote the z- and p-values corrected for multiple comparisons by the 2654 single-step method, bold values are statistically significant. 2655

2656

2657 **Diet quality effect on copepods.** The energy content of *M. littorale* was related to the stressor-2658 induced changes in diatom energy content (both stressors) and diatom community structure 2659 (atrazine only, Fig. 5.3, Table 5.3). Copepods maintained their control EFA content when feeding 2660 on diatoms from the low copper treatment, but lost half of their EFAs when feeding on diatoms 2661 grown under high copper stress (Fig. 5.3). This resulted in a positive correlation of copepod EFA 2662 content and diatom EFA content (Fig. 5.3, Table 5.3). When offered diatoms from the low and high 2663 atrazine treatments, copepods also lost half of their EFA content, which was predicted not only by 2664 the diatoms' EFA content, but also by the changes in diatom community structure (Fig. 5.3, Table 2665 5.3). Addendum IV Fig. S3 shows the EFA content per diatom biomass and per copepod individual, 2666 as well as the relative proportion of eicosapentaenoic acid (EPA) and docosahexaenoic acid (DHA). 2667 In diatoms, the essential fatty acids consisted mainly of EPA, whereas DHA was the main 2668 component in copepods (EPA:DHA ratio 4 to 6 in diatoms, 0.3 to 0.6 in copepods, Addendum IV 2669 Fig. S3). The copepod survival rate in the experimental units was between 85-100%. The EFA

- 2670 content of copepods feeding on diatoms from control conditions (65.3 ± 6.1 ng copepod-1) did not
- 2671 differ significantly from copepods at the start of the experiment (i.e. animals collected in the field,
- 2672 68.8 ± 11.8 ng copepod<sup>-1</sup>, t-test, p=0.99, Addendum IV Fig. S3).



2674

2675 Fig. 5.3: Community structure and energy content (EFA content) of diatoms exposed to stress, and energy 2676 content (EFA content) of *Microarthridion littorale* copepods after 10d of feeding on the respective diets, 2677 visualised as percent dissimilarity from the corresponding control. The dissimilarity of diatom community 2678 structure is calculated as the Bray-Curtis similarity to the control mean (see methods section). The 2679 dissimilarity of diatom and copepod EFA content is calculated as fraction percentage of the EFA 2680 concentration in treatment *i* and the mean ( $\mu$ ) EFA concentration in corresponding control c: 2681  $(EFA_i/\mu EFA_c)x100$ . Diatom communities were exposed to low (200  $\mu g/L$ ) and high (500  $\mu g/L$ ) atrazine 2682 ('Atr') and copper ('Cu') concentrations, copepods were not exposed to any of the stressors.

2683

Stressor	Quality	Model	Slope	s.e.	Т	Р	AIC	Log lik.	Validity	LR	P LR
Atrazine	EFA	1	0.059	0.008	7.38	0.0002	28.46	-11.23	Yes	0.69	0.41
		2	0.060	0.007	9.06	<0.0001	29.77	-10.89	Yes		
	Comp	1	0.045	0.008	5.88	0.0006	31.69	-12.85	Yes	0.07	0.79
		2	0.045	0.008	5.64	0.0008	33.62	-12.81	Yes		0.7 5
Copper	EFA	1	0.137	0.021	6.57	0.0003	35.21	-14.61	Yes	3.64	0.06
		2	0.133	0.014	9.17	<0.0001	33.57	-12.79	Yes		
	Comp	1	0.204	0.126	1.62	0.1486	44.82	-19.41	No	0.75	0.39
	Comp	2	0.071	0.150	0.48	0.6494	46.07	-19.03	Yes		0.57

2685

2686 Table 5.3: Results of generalized least squared models predicting copepod energy content (EFA content) 2687 from diatom community structure and energy content (EFA production). 'Stressor' denotes which diet the copepods were offered: diatom communities exposed to copper or atrazine. 'Quality' indicates the 2688 2689 predictor: diet quality in terms of diatom community composition or EFA production. 'Mod' indicates if the 2690 model was fitted without (Model 1) or with (Model 2) variance structure. 'Slope' indicates the relation 2691 between predictors and copepod energy content, i.e. the effect of diatom energy content and community 2692 structure on copepod energy content. 's.e.' is the standard error on the estimated slopes. 'T' and 'P' denote 2693 the t- and p-values, bold values are statistically significant.; 'AIC' is the Akaike information criterion, 'Log 2694 lik' the log-likelihood. 'Validity' denotes if residuals were homogeneous and normally distributed ('yes') or 2695 not ('no'). If 'no', models were refitted ('Model 2') with a variance structure allowing the residuals to change 2696 with the predictor. 'LR' is the likelihood ratio of model 1 vs. model 2, P LR the corresponding p-value. 2697

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## 2700 **5.4. Discussion**

The outcome of our experiments demonstrates that chemical stress can stimulate the contribution
of marine primary producers to sediment stabilization but reduces diet quantity and diet quality.
Selective and context-dependent stress effects on diet quality were furthermore found to lead to
concomitant knock-on effects at the consumer level.

2705 Biomass production is the most widespread functional endpoint in research investigating 2706 ecosystem functioning under anthropogenic change, and producer biomass (diet quantity) has 2707 been the emphasis of most trophic experiments (Arts and Wainmann 1999, Balvanera et al. 2006, 2708 Cardinale et al. 2011). The loss of diatom biomass induced by copper corresponds to previous 2709 findings on the toxicity of copper to marine diatoms (Stauber and Florence 1989, Cid et al. 1995, 2710 Pistocchi et al. 1997, Masmoudi et al. 2013). Surprisingly, diatom biomass was less affected by 2711 atrazine, although atrazine has been shown to reduce diatom biomass at concentrations lower 2712 than those used in our study (Bester et al. 1995, DeLorenzo et al. 1999, 2001, Magnusson et al. 2713 2008). The capacity of diatom communities to maintain their biomass under atrazine exposure was related to a change in community composition, with *C. closterium* increasing its biomass in 2714

2715 the presence of atrazine, and dominating all communities in the atrazine treatments. Atrazine 2716 blocks the electron transport chain of Photosystem II (PSII, Dorigo and Leboulanger 2001). Some 2717 diatom species can however reduce their dependency on photosynthesis and thus their sensitivity 2718 to PSII inhibitors by mixotrophic growth (i.e. the uptake of organic substrates, Debenest et al. 2719 2009; Larras et al. 2014). C. closterium is capable of mixotrophic growth, which reduces its 2720 sensitivity to herbicide stress (Vanelslander et al. 2009, Chapter 3). The presence of a mixotrophic, 2721 atrazine-tolerant species thus caused the changes in community composition and underpinned 2722 the limited loss of biomass under atrazine.

2723 Neither atrazine nor copper reduced the diatoms' contribution to sediment stabilization. On the 2724 contrary, EPS production doubled at low levels of both stressors. In our low atrazine and copper 2725 treatments, diatom EPS production was also double than expected in synthetic communities of 2726 the same community structure. The increase in EPS production thus resulted from context-2727 dependent effects rather than from changes in community structure. The secretion of EPS by 2728 microphytobenthos does not only play a vital role for ecosystem functioning through the 2729 stabilization of sediment surfaces, but also represents a physiological response of benthic diatoms 2730 to various abiotic stressors (Pistocchi et al. 1997, Decho 2000, Staats et al. 2000a, Underwood and 2731 Paterson 2003, Levy et al. 2007). At sublethal stress levels, context-dependent stress effects are 2732 thus unlikely to cause a reduction in the diatom's contribution to sediment stabilization. Selective 2733 stress can reduce diatom EPS production, by causing dominance by tolerant species producing 2734 little amounts of EPS (Mensens et al. 2015). The dominant species in our atrazine, copper and 2735 control communities however showed little difference in their EPS output, which resulted in an 2736 absence of selective stress effects on EPS production.

2737 Selective atrazine stress determined the energy content of diatom communities, which lost 60% 2738 and 75% of their essential fatty acids at low and high levels of the pesticide. The same extent of energy loss was observed in synthetic communities which reflected the community structure 2739 2740 under atrazine without actual exposure to the pesticide. Conversely, the energy content in 2741 atrazine-exposed experimental communities and unexposed synthetic communities did not differ. 2742 The energy loss under atrazine was thus caused by selective changes in community structure 2743 rather than by context-dependent atrazine effects on diatom energy content. This selective 2744 atrazine stress was underpinned by the dominant species *C. closterium*, which produced eight and 2745 three times less EFAs than the species contributing the most biomass under control conditions (N. 2746 acicularis and N. arenaria respectively, see fatty acids per species in Addendum IV Table S1). In 2747 the presence of copper, communities were dominated by the same lipid-rich species as in the 2748 control, and no such selective effects on energy content were observed. The loss of diatom energy 2749 content at high copper levels was caused by context-dependent rather than selective copper

2750 effects, which was reflected in the lower energy content in experimental compared to synthetic 2751 communities. Context-dependent effects also likely caused the further loss of EFAs at high 2752 compared to low atrazine stress, since community structure at the two herbicide levels did not 2753 differ. This loss of energy content at high levels of both stressors could be due to physiological 2754 stress effects such as an alteration of photosynthesis and thus of the carbon supply for fatty acid 2755 synthesis, an inhibition of the enzymes involved in lipid biosynthesis or an increase in the degree 2756 of fatty acid saturation, which are all reported to reduce the EFA production of microalgae under 2757 metal and pesticide stress (Böger et al. 2000, Guschina and Harwood 2006, Chia et al. 2013a).

2758 Our results however highlight selective stress as the main driver of primary producer energy 2759 content. Selective atrazine stress caused a more important loss of diatom energy content than the 2760 context-dependent effects of both stressors, at chemical concentrations where no context-2761 dependent effects on individual fatty acid content were recorded. The high and low copper and 2762 atrazine concentrations used in our experiments are respectively far above field concentrations, 2763 or have only been recorded in extreme pollution events or at chronically contaminated sites 2764 (Millward and Grant 2000, Graymore et al. 2001, Pennington et al. 2001, Lockert et al. 2006). 2765 Apart from scenarios of severe chemical pollution, pesticide and metal stress are thus unlikely to 2766 reduce the energy flow in intertidal systems by inhibiting algal EFA synthesis. Conversely, 2767 chemical pollution can cause shifts in algae community structure at stress levels lower than those 2768 used in our study (Bérard and Benninghoff 2001, Debenest et al. 2010). The impact of such 2769 selective chemical stress on ecosystem functioning will largely depend on the functional 2770 importance of the most stress-tolerant species (Larsen et al. 2005, Mensens et al. 2015, Radchuk 2771 et al. 2016). Not only pollution, but also long-term changes in environmental variables are causing 2772 considerable shifts in microalgae community structure (Pomati et al. 2012, Litchman et al. 2015). 2773 Due to the pronounced differences in lipid profiles within and among algal classes (Dunstan et al. 2774 1993, Taipale et al. 2013, Guo et al. 2016), these changes in community structure rather than 2775 context-dependent changes in algal energy content represent the potentially stronger driver of 2776 trophic energy flow under anthropogenic change.

2777 The energy content of the copepod *M. littorale* closely tracked that of its diets. Selective and 2778 context-dependent stress effects on diatom energy content resulted in a concomitant loss in the 2779 energy content of their main copepod grazer, confirming algal EFAs as a key component of diet 2780 quality which is directly linked to trophic energy transfer. For diatoms exposed to atrazine, the 2781 response of *M. littorale* was predicted by both its diet's energy content and community structure. 2782 Energy content in terms of fatty acids is the most common measure of diet quality (Jodice et al. 2006, Österblom et al. 2008, Taipale et al. 2013, Guo et al. 2016), but requires extensive analyses 2783 2784 of a diet's biochemical composition. For stressors causing selective shifts in community structure,

effects on trophic energy transfer can thus more straightforwardly be estimated from the speciesdominating under stress, given that the nutritional value of these species is known.

2787 However, our results have to be treated carefully due to the limited number of stress levels and 2788 replicates. To obtain a more robust correlation of the consumer response to stressor-induced 2789 changes in diet quality, models should be fitted to gradients containing more than three 2790 concentrations per stressor and more than three replicates per treatment. Also, offering M. 2791 *littorale* preserved rather than live diatom food might have influenced food uptake. Freeze-drying 2792 does not alter the biochemical composition within diatoms cells, but modifies the exterior of 2793 diatom cells through the loss of exudates (e.g. EPS) or bacteria associated to the diatom frustule, 2794 which can affect the ingestion of diatoms by harpacticoid copepods (Cnudde et al. 2011). Feeding 2795 *M. littorale* live diatom cultures under unstressed conditions would however have resulted in a 2796 dissimilar diatom community structure than that induced by the stressors: atrazine and copper 2797 did not eliminate any of the diatom species, but caused alterations of community evenness, which 2798 could not be maintained when diatoms were grown in the absence of the stressors.

2799 The DHA:EPA ratio of *M. littorale* was higher than in the diatom communities, which corresponds 2800 to previous findings on the relative concentrations of both essential fatty acids in copepods and 2801 their algal diets (De Troch et al. 2012a, Arndt and Sommer 2014). Diatoms are characterised by a 2802 high EPA content (Taipale et al. 2013, Guo et al. 2016), but DHA appears to be the most important 2803 fatty acid for copepods (Taipale et al. 2013). Planktonic primary consumers such as cladocerans 2804 or calanoid copepods directly depend on the DHA taken up from their diet (Bell et al. 2007, Bell 2805 and Tocher 2009, De Troch et al. 2012a), but several harpacticoid copepod species are able to 2806 bioconvert EPA to the longer chain DHA, a capacity which has notably been demonstrated in M. 2807 *littorale* (De Troch et al. 2012a). While the total EFA content of *M. littorale* reflected that of its 2808 different diatom diets, this capacity to convert EPA to DHA likely enabled *M. littorale* to maintain 2809 high relative levels of DHA.

2810 Since their EFA content ranks among the highest of all algae classes, diatoms are regarded as high-2811 quality food source and crucial link for the energy flow at the basis of aquatic food webs (Brett 2812 and Müller-Navarra 1997, Taipale et al. 2013, Guo et al. 2016). Here, diatom diet quality 2813 responded more sensitively to chemical stress than the diatom's contribution to habitat structure 2814 (sediment stabilization) and diet quantity. The selective and context-dependent stress effects on 2815 diatom diet quality were caused by the large interspecific differences in energy content and the 2816 loss of diatom energy content under stress. Indeed, the energy content of benthic diatoms shows 2817 more interspecific variation and is more affected by chemical pollutants than their contribution 2818 to diet quantity or sediment stabilization (Mensens et al. 2015, Chapter 3).

2819 Losses in diet quality occurred at copper concentrations which were close to the LC50 value of M. 2820 *littorale* (Addendum V). Next to indirect effects on copepod energy content through changes in 2821 diet quality, high metal stress can thus be expected to reduce copepod abundance through direct 2822 toxic effects. Conversely, atrazine changes microalgal community structure at concentrations 2823 lower than the 200 µg/L used in our study (Bérard and Benninghoff 2001, Debenest et al. 2010), 2824 whereas direct acute effects of atrazine on copepods and other marine invertebrates mostly occur 2825 at >1 mg/L atrazine (Hall et al. 1995, Bejarano and Chandler 2003). Herbicides can thus first affect 2826 marine primary consumers indirectly through selective changes in the structure of their diet 2827 rather than through direct effects on the consumers themselves. The loss of diet quality did not 2828 result in increased harpacticoid mortality. Low diet quality rarely causes acute copepod mortality, 2829 but reduces copepod energy content and, on a longer term, growth rate and reproduction (Müller-2830 Navarra 1995, Müller-Navarra et al. 2000, Arendt et al. 2005, Gonçalves et al. 2011). The energy 2831 content of copepods is crucial for their main consumers, especially larval fish whose development 2832 can depend on the EFAs taken up from their copepod prey (Sargent et al. 1995, Payne et al. 1998). 2833 Rather than acutely eliminating consumers, the main risk of losses in producer diet quality thus 2834 consists in reducing the energy transfer at the plant-animal interface, which is a key limiting factor 2835 for the functioning of marine ecosystems (Brett and Müller-Navarra 1997, De Troch et al. 2012b). 2836 Nonetheless, the importance of algal diet quality, as distinct from just diet quantity, is rarely 2837 highlighted in research on food web functioning (Guo et al. 2016). Due to its sensitive response to 2838 selective stress, algal diet quality in terms of EFA production and community structure provides 2839 a powerful approach to integrate our understanding of coastal ecosystem functioning under 2840 anthropogenic change.

2841 In conclusion we infer three statements from our results. First, chemical stress differentially 2842 affects the contribution of marine primary producers to diet quantity, sediment stabilization and 2843 diet quality, with diet quality being most sensitive. Second, the loss of diet quality is underpinned 2844 by selective and context-dependent stress effects. Selective stress causes a more important loss of 2845 diet quality at lower stress levels than context-dependent stress effects, and thus represents the 2846 main risk for energy flow at the basis of marine food webs. Third, changes in producer diet quality 2847 closely predict consumer energy content. The integration of diet quality within traditional studies 2848 of diet quantity is recommended to assess energy flow in marine food webs under anthropogenic 2849 change.

2850

## 2851 6. General discussion

2852 Since the first BEF experiments were conducted in the 1990s (Naeem et al. 1994, Tilman and 2853 Downing 1994, Tilman et al. 1996, Hector et al. 1999), nearly a thousand studies have examined 2854 the functional role of biodiversity in terrestrial, freshwater and marine ecosystems (Covich et al. 2855 2004, Cardinale 2011, Cardinale et al. 2012, Gamfeldt et al. 2015). After initially controversial 2856 discussions (van der Heijden et al. 1999, Wardle 1999, Hector et al. 2000, Huston et al. 2000), 2857 consensus is now emerging that declines in biodiversity have negative consequences for 2858 ecosystem functioning (for quantitative data syntheses see Worm et al. 2006, Balvanera et al. 2859 2006, Cardinale et al. 2006, 2011, 2012, Stachowicz et al. 2007, Finkel et al. 2010, Quijas et al. 2860 2010, Hooper et al. 2012, Tilman et al. 2014, Gamfeldt et al. 2015, Hautier et al. 2015). In addition, 2861 BEF research has generated new ecological and mathematical theory (Tilman et al. 1997, Loreau 2862 and Hector 2001, Pacala and Kinzig 2002, Fox 2005), provided decision support tools for 2863 policymakers (Naidoo and Ricketts 2006, Kareiva et al. 2011, InVEST 2016), and led to the 2864 establishment of intergovernmental initiatives to preserve global biodiversity (CBD 2010, 2865 Larigauderie and Moonev 2010, IPBES 2016).

2866 Despite this progress, early (Hooper et al. 2005) and recent (Cardinale et al. 2012, Tilman et al. 2867 2014, De Laender et al. 2016) BEF consensus papers continue to highlight the need to include the 2868 environmental drivers of biodiversity loss into the next generation of BEF experiments. 2869 Contrasting BEF relations under stress with the classic BEF protocol of random species loss, and 2870 the identification of response and effect traits capable of predicting biodiversity and ecosystem 2871 functioning under stress, have been identified among the most urgent need to explore the 2872 functional consequences of realistic scenarios of biodiversity change (Suding et al. 2008, Naeem 2873 2008, Cardinale et al. 2012, Tilman et al. 2014, Wardle 2016).

2874 This dissertation, together with other recent efforts to include stressors into BEF theory (see 2875 further), has shown that stress can profoundly alter the biodiversity effect on ecosystem 2876 functioning. Diversity loss under stress is not **random**, but **selective** according to the species' 2877 stress tolerance, and the functional contribution of the most tolerant species plays a key role for 2878 functioning under stress. BEF relations in diatom communities under atrazine were steeper than 2879 predicted by the classic random design, since the herbicide inhibited the growth of the species 2880 (*Nitzschia* sp.) contributing most to energy content and sediment stabilization, and communities 2881 were dominated by a tolerant species (*N. arenaria*) contributing little to both processes (Chapter 2882 2). Theory predicts that under stress, the abundance and functional contribution of species can be 2883 predicted from their biological traits. The numerical and functional response of diatoms to copper 2884 stress was related to the same set of response and effect traits (cell size and surface-to-volume

2885 ratio), whilst the diatoms' capacity for mixotrophic growth predicted the numerical but not the 2886 functional stress response to atrazine (Chapter 3). **Complementarity**, the main statistical effect 2887 used to explain the better performance of species in community, has rarely been quantified under 2888 stress and it is unclear what **mechanisms** underpin potential changes in complementarity. 2889 Complementarity in diatom communities increased under atrazine and copper stress, driven by 2890 facilitation (an increased release of extracellular polymers). Complementarity benefited the 2891 growth of mixotrophic species (e.g. C. closterium) under atrazine, and of copper-sensitive species 2892 with low biomass yields (e.g. A. lineolata) under metal stress, reducing the net diversity effect on 2893 biomass yield in the latter case (Chapter 4). There is little evidence whether stressors drive 2894 functioning through selective effects (changes in community structure) or context-dependent 2895 effects (changes in the species' functional contribution), and if both types of stress knock on 2896 across **trophic levels**. Selective atrazine effects, leading to dominance by a mixotrophic but lipidpoor species (*C. closterium*), caused a 60% loss of diatom energy content at low atrazine levels. 2897 Context-dependent stress effects only affected diatom functioning at high atrazine and copper 2898 2899 levels. Both types of stress affected the energy transfer to the next trophic level, causing a 2900 concomitant loss in the energy content of their main copepod consumer (*M. littorale,* Chapter 5).

2901 Based on these findings, three main effects that potentially drive functioning in stressed 2902 communities can be put forward. This discussion will first focus on selective stress effects, i.e. 2903 shifts in community structure causing stressed communities to be dominated by species with an 2904 above- or below-average functional contribution. Subsequently, context-dependent stress effects 2905 on the species' functional contribution will be addressed, which can be mediated by **physiological** 2906 stress effects and changes in complementarity, i.e. differential interspecific interactions in 2907 stressed communities. These three effects are not inherently exclusive, could not always be 2908 separated, and will in most cases simultaneously determine the impact of stressor-induced 2909 biodiversity changes on ecosystem functioning. Nevertheless, the most important stressor-2910 induced changes in the contribution of diatom communities to ecosystem functioning, including 2911 concomitant effects at the consumer level, can be interpreted with regard to these three effects. 2912 Each effect is first discussed (sections 6.1 to 6.3) and then examined with regard to its impact on 2913 three ecosystem processes (biomass production, sediment stabilization and energy content<sup>1</sup>, 2914 section 6.4), before presenting future challenges for BEF research which arise from the outcome 2915 of this work (section 6.5).

<sup>&</sup>lt;sup>1</sup> Energy content was quantified as polyunsaturated fatty acids (PUFAs, fatty acid molecules with more than one double bond) in Chapters 2 and 3, and as essential fatty acids (EFAs, the two most important PUFAs for animal physiology: EPA and DHA) in Chapter 5. For the sake of simplicity, this chapter only uses the terms 'energy content' and 'fatty acids'.

#### 2916 **6.1. Selective stress**

2917 Selective stress (Fig. 6.1.1) can be defined as a change in environmental conditions which affects 2918 ecosystem functioning by selectively targeting species with a high or low functional contribution 2919 (Chapter 1.4.2). Random species loss is the classic protocol in BEF research (Naeem 2008, 2920 Cardinale et al. 2012, Wardle 2016). Theory and simulations however predict stress to cause non-2921 random scenarios of biodiversity loss, where species are targeted selectively according to their 2922 stress tolerance, and the functional contribution of the most tolerant species plays a key role for 2923 the overall functioning in stressed communities (Solan et al. 2004, Bunker et al. 2005, Cardinale 2924 et al. 2012).

2925 Here, stressor-induced changes in diatom community structure had little effect on the biomass 2926 production of stressed diatom communities, which depended on complementarity (see 6.3) rather 2927 than on the biomass production of the dominant species (Chapter 4). Selective atrazine effects 2928 reduced the contribution of stressed diatom communities to sediment stabilization, by causing 2929 dominance of an atrazine-tolerant species (N. arenaria) which however produced little EPS 2930 (Chapter 2). When stress-sensitive and stress-tolerant species did not differ in their functional 2931 contribution, selective changes in community structure however had no effect on the contribution 2932 of stressed diatom communities to sediment stabilization (Chapter 5). The strongest selective 2933 stress effects were recorded in terms of diatom energy content. Selective atrazine effects inhibited 2934 the growth of a lipid-rich species (Nitzschia sp.), causing a disproportionate reduction in the 2935 energy content of diatom communities exposed to the herbicide, which was not predicted by the 2936 classic design of random species removal (Chapter 2). Similarly, atrazine induced a shift in diatom 2937 community structure from dominance by a lipid-rich species (*N. acicularis*) under control 2938 conditions towards dominance by a highly mixotrophic, but lipid-poor species (*C. closterium*) 2939 under atrazine (Chapter 5). This selective shift in community structure caused a 60% loss in the 2940 energy content of diatom communities under atrazine, which furthermore knocked on to the next 2941 trophic level and reduced copepod energy content by half, despite invariant diatom biomass and 2942 in the absence of any context-dependent stress effects. Selective atrazine stress thus represented 2943 the main risk for energy flow at the producer-consumer interface. Conversely, no selective copper 2944 effects on diatom energy content were detected, since exposure to the metal did not change 2945 diatom community structure (Chapter 5).

Moreover, BEF research has been sparked by concerns about the loss of species and, with few exceptions (e.g. Balvanera et al. 2005, Wittebolle et al. 2009), the majority of BEF experiments have manipulated diversity in terms of species richness, holding evenness constant across richness treatments (Hillebrand et al. 2008, Hillebrand and Matthiessen 2009, Zhang et al. 2012). However, critics often assert that anthropogenic stress is more likely to alter evenness than richness, with changes in evenness potentially affecting ecosystem functioning before species are driven to extinction (Chapin et al. 2000, Balvanera et al. 2005, Hillebrand et al. 2008). Here, atrazine and copper consistently altered evenness, but did not cause any species loss, highlighting the need to investigate diversity loss not only in terms of species extinctions, but also altered community structure. More importantly, when changes in evenness involved the dominance of species with low functional contribution, loss of function was more important than could be predicted from randomly composed communities (Chapter 2).

- 2958 Selective biodiversity loss can in theory cause more rapid or slower declines in ecosystem 2959 functioning compared to random loss scenarios (Ives and Cardinale 2004, Solan et al. 2004, 2960 Bunker et al. 2005, Gross and Cardinale 2005), and experimental evidence suggests the functional 2961 impact of selective stress depends on the stress tolerance of those species contributing most to 2962 functioning (Larsen et al. 2005, McIntyre et al. 2007, Ball et al. 2008, Bracken et al. 2008, Davies 2963 et al. 2011, Radchuk et al. 2016). For instance, diversity loss caused by accelerated wave exposure 2964 and overfishing reduced nutrient cycling in macroalgae and fish communities respectively to a 2965 larger extent than expected from random diversity loss (McIntyre et al. 2007, Bracken et al. 2008). 2966 In both cases, wave stress and fishermen targeted the species contributing most to nutrient cycling. In the terrestrial environment, habitat loss caused a disproportionate loss of pollination, 2967 2968 because large-bodied insects contributing most to pollination were most sensitive to this type of 2969 stress (Larsen et al. 2005). Conversely, insecticide stress had no effect on ecosystem processes in 2970 an aquatic food web, since the insecticide only targeted species contributing little to functioning, 2971 and the functionally most important species could maintain their functional contribution under 2972 stress (Radchuk et al. 2016).
- Suding *et al.* (2008) and Hillebrand and Matthiessen (2009) highlighted the need for BEF research to consider the functional consequences of selective biodiversity loss, by proposing a trait-based framework which connects species traits that influence the stress tolerance ('response traits') to species traits that influence ecosystem functioning ('effect traits'). Although experimental tests of this framework are still rare (but see Heuner et al. 2015, Lennon and Lehmkuhl 2016), relating response and effect traits has been identified as a key issue to estimate ecosystem functioning under stress (Cardinale et al. 2012, Wardle 2016).

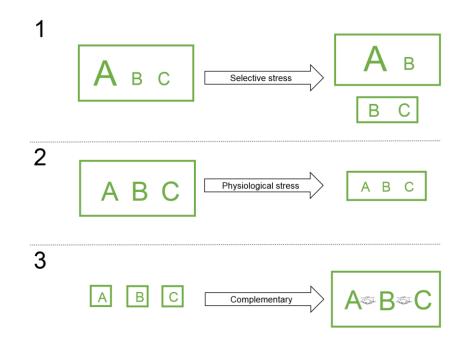
Here, the response-effect trait relationship was different depending on stressor type. The capacity of diatom species to withstand copper effects on density ('numerical response') and on functional contribution ('functional response', in terms of photosynthetic efficiency, energy content and sediment stabilization) was predicted by the same set of morphological traits (cell size and surface to volume ratio). Species which were numerically tolerant to copper contributed most to functioning under unstressed conditions, and performed best at maintaining functioning when exposed to the metal, indicating that tolerant species might be able to sustain functioning under copper stress. A physiological trait (mixotrophy) predicted the diatoms' numerical response to atrazine, but not their functional response. Numerically tolerant diatoms did not contribute most to functioning, and the numerical and functional response to atrazine were not correlated, indicating a limited capacity of tolerant species to maintain functioning under atrazine (Chapter 3).

2992 The diatoms' numerical stress response to atrazine and copper corresponded to the community 2993 structure observed under both stressors: copper-exposed diatom communities were dominated 2994 by Entomoneis sp. and N. acicularis (Chapters 4 & 5) which were among the numerically and 2995 functionally most tolerant species to the metal. Consequently, the functioning of diatom 2996 communities was only affected at high (500 µg/L) copper concentrations. Atrazine-exposed 2997 diatom communities were dominated by two species which were numerically tolerant to the 2998 herbicide (*N. arenaria* and *C. closterium*). These species contributed seven times less to sediment 2999 stabilization and respectively fifteen and fourty times less to energy content than the atrazine-3000 sensitive species which had dominated communities under unstressed conditions (*Nitzschia sp.* 3001 and *N. acicularis*, Chapter 2 and 5), causing the negative selective atrazine effects on diatom 3002 functioning (see above). Entomoneis sp. and C. closterium were the most copper- and atrazine-3003 tolerant species among a set of 17 diatom species, and were characterized by the largest cell size 3004 and lowest surface-to-volume-ratio, and by the highest capacity for mixotrophic growth 3005 respectively (Chapter 3).

3006 Whilst the above examples confirm the potential of traits to predict changes in community 3007 structure and their effect on functioning under stress (Hillebrand and Matthiessen 2009), this 3008 trait-based approach could not consistently predict community structure under stress: For 3009 instance, N. arenaria reached high relative abundances in atrazine-exposed communities (Chapter 3010 2), despite having no capacity for mixotrophic growth (Chapter 3). Similarly, the good 3011 performance of *N. acicularis* in copper-exposed communities (Chapter 5) was not predictable 3012 from its traits, since this species was neither characterized by a large cell size nor by a low surface 3013 to volume ratio (Chapter 3). Selective stress effects on diatom community structure and 3014 functioning were thus consistently predicted by the species' tolerance (EC50 values), but not 3015 always by the species' traits, arguably because this work did not include other traits relevant to 3016 microalgal stress tolerance. Such traits involve metal-binding metallothioneins and 3017 phytochelatins, or the structure of the quinone binding site in photosystem II, which determine 3018 algal tolerance to copper and atrazine respectively (Galloway and Mets 1984, Gekeler et al. 1988, 3019 Robinson 1989, Fernandes and Henriques 1991, Galgani et al. 1999). Furthermore, stress

3020 tolerance in monoculture, which was predicted by response traits, could be decoupled from stress 3021 tolerance in community when species benefited differentially from complementarity. For 3022 example, in the diatom communities in Chapter 4, A. lineolata was the least copper-tolerant 3023 species in monoculture, which could be expected based on its response traits (smallest cell size, 3024 largest surface to volume ratio of all component species). A. lineolata however benefited most 3025 from an increase in complementarity and thus performed well in communities exposed to the 3026 metal (but see 6.3), thus limiting the capacity of response traits to predict community structure 3027 under stress.

3028 Nevertheless, the identification of key morphological and physiological traits determining, in part, 3029 the structure and functioning of primary producer communities under stress, highlighted the 3030 value of trait-based approaches to predict the consequences of selective stress for ecosystem 3031 functioning. In ecotoxicology, approaches relating traits to stress tolerance have been tested 3032 (Baird and Van den Brink 2007, Rubach et al. 2012, Larras et al. 2014) and trait-based bioindicator 3033 tools are being used to link the pesticide load in European waters to the structure of aquatic 3034 communities (Liess and Von Der Ohe 2005, Beketov et al. 2009). In BEF research, the theoretical 3035 framework for the incorporation of trait-based approaches has been set (Suding et al. 2008, 3036 Hillebrand and Matthiessen 2009, Litchman et al. 2015). Methodological advances now allow for 3037 time-efficient quantifications of traits at the individual level (Fontana et al. 2014), and recent 3038 experimental approaches have used response and effect traits to understand the structure and 3039 functioning of communities under stress (Koide et al. 2014, Heuner et al. 2015, Zwart et al. 2015, 3040 Lennon and Lehmkuhl 2016). These theoretical and methodological advances should allow to 3041 progressively complement the classic design of random species loss with trait-based approaches 3042 that more realistically reflect community structure and ecosystem functioning under 3043 anthropogenic change.



3046 Fig. 6.1: Scheme indicating three effects which potentially drive functioning in stressed communities. For 3047 the sake of simplicity, this scheme uses communities with only three species (A, B and C). The font size of A, 3048 B and C indicates the species' functional contribution. Communities are marked by green boxes, with the 3049 size of the box marking the community's functional contribution. Panel 1 shows selective stress (for 3050 details see Chapter 1.4.2), which occurs when the species contributing most to functioning are also least 3051 stress-tolerant. The example community in Panel 1 includes a species with a high functional contribution 3052 (species A) and two species with a low functional contribution (species B and C). Selective stress has limited 3053 effect on community functioning, if the stressor does not remove species A, which contributes most to 3054 functioning (upper community on the right-hand side of panel 1). Selective stress disproportionately 3055 reduces community functioning, if the stressor removes species A (lower community on the right-hand side 3056 of panel 1). **Panel 2** shows **physiological stress** (for details see Chapter 1.4.3), which affects ecosystem 3057 functioning when stress alters the species' functional contribution. For the sake of simplicity, Panel 2 3058 indicates a community in which the species, A, B and C contribute equally to functioning, with each species' 3059 functional contribution altered to the same extent by stress. Panel 3 shows complementarity (for details 3060 see Chapter 1.6), which occurs when species contribute more to functioning when grown in community 3061 than in monoculture. Facilitation and niche partitioning are the main mechanisms driving complementarity. 3062 In contrast to selective and physiological stress, complementarity is not a 'stress effect', and can operate 3063 under stressed and unstressed conditions. The example in Panel 3 shows three species A, B and C with a 3064 low functional contribution when being exposed to stress in monoculture, and an increased functional 3065 contribution when being exposed to stress in community due to facilitative interactions.

3066

## 3067 6.2. Physiological stress

Physiological stress can be defined as stress which affects ecosystem functioning by directly 3068 3069 altering the species' functional contribution (Fig. 6.1.2). Direct stress exposure to anthropogenic 3070 stressors is rarely included into the design of BEF experiments (McMahon et al. 2012, Steudel et 3071 al. 2012), which remove species to mimic stressed ecosystems, but assume that the functional 3072 contribution of the species in these low-diversity-communities remains unchanged. However, 3073 stressors that cause species loss can impose a physiological cost on the surviving species, e.g. 3074 acclimation by allocating resources from growth to survival pathways, which can affect the 3075 contribution of stressed communities to ecosystem functioning (Schimel et al. 2007, Hawlena and 3076 Schmitz 2010). This has resulted in calls to consider sublethal physiological effects on the species'
3077 functional contribution, when estimating ecosystem functioning under stress (Relyea and
3078 Hoverman 2006).

3079 In monoculture, atrazine and copper reduced diatom growth and thus biomass production 3080 (Chapters 3 & 4). The main physiological mechanisms through which atrazine and copper affect 3081 diatom growth comprise an inhibition of electron transport in photosystem II (atrazine, Dorigo 3082 and Leboulanger 2001) and the damage of cell membranes, thylakoids and chloropolasts (copper, 3083 Knauert and Knauer 2008, for more details see Chapter 1.8.1). Conversely, most diatom species 3084 increased their contribution to sediment stabilization by producing more EPS at low atrazine and 3085 copper levels (Chapter 3). At high levels of both stressors diatoms could not maintain their EPS 3086 production (Chapter 3). As well as playing an important functional role in sediment stabilization 3087 (De Brouwer et al. 2000, Decho 2000), the excretion of EPS also represents a physiological stress 3088 response of benthic diatoms to metal, nutrient and salinity stress (Pistocchi et al. 1997, Liu and 3089 Buskey 2000, Staats et al. 2000a, Levy et al. 2007). However, diatoms cannot maintain their EPS 3090 production when photosynthesis is inhibited (e.g. by herbicides, Staats et al. 2000b), which can 3091 explain the loss of EPS at high stress levels. Diatom energy content was consistently reduced by 3092 both stressors, with most diatom species producing less fatty acids at stress levels where their 3093 growth was still unaffected (Chapter 3). Physiological pesticide and metal effects that can lead to 3094 such losses in algal energy content include a reduced carbon supply for fatty acid synthesis (when 3095 photosynthesis is inhibited), the conversion of unsaturated to saturated fatty acids, or the 3096 inhibition of enzymes involved in fatty acid synthesis (Böger et al. 2000, Guschina and Harwood 3097 2006, Chia et al. 2013a, 2013b).

To what extent did these physiological effects impact the contribution of stressed diatom communities to ecosystem functioning? Linking physiological responses to environmental stress to processes at the ecosystem level has been a major theme in plant ecology (Grime 1977, Chapin et al. 2002). The few BEF experiments conducted under stressful conditions could not disentangle whether changes in ecosystem functioning were mediated by biodiversity loss, or by effects of environmental disturbance on the species' functional contribution (Steudel et al. 2012, Radchuk et al. 2016, but see Fox and Harpole 2008, Baert et al. 2016).

Stressors can affect ecosystem functioning through selective biodiversity loss (e.g. by targeting
species with a high functional contribution, see 6.1), or through 'context-dependent effects', i.e.
changes in the species' functional contribution (Fox 2006, Fox and Harpole 2008, Tylianakis et al.
2008, Hiddink et al. 2009). These changes in functional contribution can arise from direct effects
of environmental drivers (Fox and Harpole 2008), i.e. the here-described physiological effects, as

well as from species interactions (e.g. a change in complementarity, Fox 2006, Fox and Harpole2008)

3112 Here, selective and context-dependent stress effects could be disentangled by contrasting 3113 functioning (energy content and sediment stabilization) in stressed diatom communities with that 3114 expected from unstressed monocultures (Chapters 2 & 5). The increased contribution to sediment 3115 stabilization (EPS production) of diatom communities under copper and atrazine was driven by 3116 positive context-dependent effects (Chapter 5). Since physiological effects of both stressors also 3117 increased the EPS production of most diatom species in monoculture (see above), it can be 3118 assumed that the increased EPS production of stressed diatom communities represented a direct 3119 physiological stress response. Nevertheless, the extent to which species interactions contributed 3120 to this increased EPS production under stress could not be quantified. No selective stress effects 3121 on EPS production were recorded when species did not differ in their functional contribution 3122 (Chapter 5). When species however differed in their functional contribution and selective stress 3123 caused dominance by an unproductive species (*N. arenaria*, Chapter 2), the resulting decline in 3124 EPS production could not be compensated by any context-dependent effects.

3125 Context-dependent stress effects generally reduced diatom energy content, with the fatty acid 3126 production of most species being reduced at stress levels which did not affect other processes. In 3127 Chapter 5, this loss of diatom energy content also knocked on to the consumer level and caused a 3128 concomitant loss in copepod energy content. However, compared to selective stress effects, 3129 context-dependent effects were of minor importance for diatom energy content. By promoting 3130 dominance of tolerant but lipid-poor species, selective atrazine effects caused a more important 3131 loss of diatom energy content than the context dependent effects of both stressors, at chemical 3132 concentrations where no context dependent effects on diatom energy content were recorded. At 3133 atrazine levels which, on average, reduced diatom fatty acid content by half (150  $\mu$ g/L, see EC50 3134 values in Chapter 3), the herbicide caused a shift in diatom community structure involving 3135 dominance by tolerant species (*C. closterium* and *N. arenaria*) producing fifteen and fourty times 3136 less fatty acids than those dominating under unstressed conditions (*Nitzschia* sp., *N. acicularis*).

When species showed large differences in their functional contribution, selective stress effects via changes in community structure were thus a more potent threat to ecosystem functioning than stress effects on the species' functional contribution. Previous work also found that whilst physiological stress effects commonly affect ecosystem processes on the short term, selective changes in community structure can have strong long-term effects on functioning, which prevail after the stressful conditions (Schimel et al. 2007). Moreover, physiological stress effects on the functional contribution of diatoms in monoculture could, in part, be compensated by positivespecies interactions in stressed communities, as will be discussed in the next section.

## 3145 **6.3. Complementarity**

3146 Complementarity effects occur when species contribute more to functioning in community than 3147 in monoculture (Fig. 6.1.3), and result from ecological mechanisms such as niche partitioning or 3148 facilitative interactions between species (Loreau and Hector 2001, Fox 2005). Experimental 3149 evidence linking these mechanisms to complementarity effects is however still scarce, and the 3150 extent to which these mechanisms broadly contribute to ecosystem functioning has yet to be 3151 confirmed (Vanelslander et al. 2009, Carroll et al. 2011, Cardinale et al. 2011, 2012). Two-way 3152 complementarity occurs when niche partitioning or facilitative interactions benefit all species, 3153 whilst one-way complementarity occurs if these mechanisms are not vice-versa and favour 3154 species with a high or low functional contribution (Fox 2005, Hector et al. 2009).

3155 As such, complementarity is not a 'stress effect', but operates under unstressed conditions, and 3156 has been in the focus of major debates in BEF research (Huston 1997, Huston et al. 2000, Cardinale 3157 et al. 2006, Fargione et al. 2007, Hodapp et al. 2016). Most of these debates have centred on 3158 whether biodiversity effects on ecosystem functioning are driven by complementarity effects, or 3159 by the competitive dominance of single, highly productive species (dominance effects, but see 3160 Chapter 1.6 and Loreau and Hector 2001, Fox 2005). Over the last decade, BEF research has shown 3161 that complementarity and dominance effects are not inherently exclusive and jointly control 3162 ecosystem functioning (Hector et al. 2009, Cardinale et al. 2012, Hodapp et al. 2016). On average 3163 over many ecosystems, each effect contributes roughly 50% of the biodiversity effect on 3164 functioning, whilst complementarity appears to be the main effect driving BEF relations in aquatic 3165 ecosystems (Cardinale et al. 2011, 2012).

3166 There is still little evidence on how complementarity is affected by anthropogenic stress (but see 3167 Wang et al. 2013, Baert et al. 2016). Potential changes of complementarity are however crucial for 3168 determining how stress will alter species' contribution to ecosystem functioning. If 3169 complementarity is not affected by stress, i.e. species interactions are the same under stressed 3170 and unstressed conditions, stress should affect the BEF relation only through selective 3171 biodiversity loss and physiological effects. The biodiversity effect on functioning would thus 3172 depend on the functional contribution of the most stress-tolerant (and thus dominant) species 3173 (Baert et al. 2016). If, however complementarity becomes more positive under stress, e.g. through 3174 an increase in facilitative interactions, adverse selective and physiological stress effects on 3175 functioning could be compensated. This potential for functional compensation among stressed 3176 communities has thus been a major subject of discussion BEF research (Fox 2006, Duffy et al.

2007, Fox and Harpole 2008) and stress ecology (Bertness and Callaway 1994, Maestre et al. 2009,
2010, He et al. 2013).

3179 Here, diatoms produced more biomass when grown in community than in monoculture. This 3180 positive biodiversity effect was caused by complementarity, which increased under stress 3181 (Chapter 4). Atrazine and copper reduced diatom biomass in monoculture (see 6.2), but this loss 3182 of biomass could partially be compensated by an increase in complementarity when diatoms were 3183 exposed to the same stress levels in community. Facilitative mechanisms can drive 3184 complementarity effects in unstressed conditions, for example in legumes and insects (Cardinale 3185 et al. 2002, Temperton et al. 2007). In microalgae, the release of organic exudates can act as a facilitative mechanism driving complementarity effects (Vanelslander et al. 2009). At present, 3186 3187 there is however no ecophysiological evidence linking changes in complementarity under stress 3188 to changing facilitative mechanisms. Under atrazine and copper stress, diatom communities 3189 increased their EPS production which, in part, predicted increases in complementarity under 3190 stress (Chapter 4). This role of EPS as a 'sleeping' facilitative mechanism, which when stimulated 3191 under stress predicted changes in complementarity highlights the two roles of microalgal EPS 3192 production: first, as an ecosystem process *per se*, promoting sediment stabilization (De Brouwer 3193 et al. 2000, Decho 2000, Gerbersdorf et al. 2009a). Second, as a physiological stress response 3194 which not only operates in monoculture (Pistocchi et al. 1997, Staats et al. 2000a), but also acts as 3195 driver of complementarity effects which helps to maintain biomass production under stress.

3196 Moreover, complementarity under stress was largely one-way, and benefited species depending 3197 on their properties: under atrazine and copper respectively, complementarity favoured the 3198 growth of mixotrophic (C. closterium, N. acicularis, Nitzschia sp.) and copper-sensitive species (A. 3199 *lineolata*, *N. digitoradiata*, *Gyrosigma* sp.). To date, the few BEF experiments which partitioned 3200 two-way and one-way complementarity found only minor contributions of one-way 3201 complementarity to the net biodiversity effect on ecosystem functioning (Hector et al. 2009, 3202 Fernandes et al. 2011, Long et al. 2013, Stachova et al. 2013, Siebenkäs et al. 2016). Here, copper-3203 sensitive species were characterised by low biomass yields, causing a negative one-way 3204 complementarity under copper stress, which largely offset positive two-way complementarity 3205 effects and thus limited the biodiversity effect on diatom biomass production. The release of 3206 extracellular polymers as a facilitative mechanism thus predicted changes in complementarity in 3207 stressed diatom communities, but was not necessarily beneficial for ecosystem functioning under 3208 stress when complementarity benefited species with a low functional contribution. In total 3209 however, complementarity was consistently positive, increased under stress when one-way 3210 complementarity did not favour species with a low functional contribution, and thus represented 3211 the key driver of diatom biomass production under stress.

## 3212 6.4. Multifunctionality

3213 The large majority of BEF research has focused on one ecosystem process at a time, most 3214 commonly the biomass production of primary producers (Balvanera et al. 2006, Jiang et al. 2008, 3215 Cardinale et al. 2011). This focus on biomass is understandable, given the importance of biomass 3216 production for e.g. food and material production or carbon sequestration, which represent 3217 important ecosystem services that human society heavily relies on (Jiang et al. 2008, Cardinale et 3218 al. 2012). Here, however, different species contributed to different ecosystem processes (see 6.1-3219 6.3). These results correspond to a growing body of BEF studies which found that sustaining 3220 multiple rather than single ecosystem processes requires a greater number of species, and 3221 suggested that more biodiversity is needed to maintain the 'multifunctionality' of ecosystems 3222 (Isbell et al. 2011, Maestre et al. 2012, Pasari et al. 2013, Lefcheck et al. 2015, Soliveres et al. 2016).

3223 If the covariation among diversity effects driving different ecosystem processes is strong, e.g. if 3224 multiple processes are regulated by either complementarity or the presence of one dominant 3225 species, a subset of diversity should be sufficient to sustain multifunctionality (Cardinale et al. 3226 2011). This principle also applies to the maintenance of multifunctionality under stress. If the 3227 diversity-multifunctionality relation is regulated by similar stress effects, e.g. by an increase in 3228 complementarity or by selective stress effects on the dominant species, focussing on one of these 3229 effects allows to predict multiple ecosystem processes under stress. If different ecosystem 3230 processes are driven by different stress effects, the maintenance of a particular ecosystem process 3231 under stress might reduce other processes, and multifunctionality imposes a tradeoff for the 3232 conservation of ecosystem functioning under stress. Here, for example, a facilitation-driven 3233 increase in complementarity stimulated the growth of a highly mixotrophic but lipid-poor species 3234 (C. closterium). This allowed to maintain diatom biomass under herbicide stress, but came at the 3235 cost of a negative selective stress effect on diatom energy content.

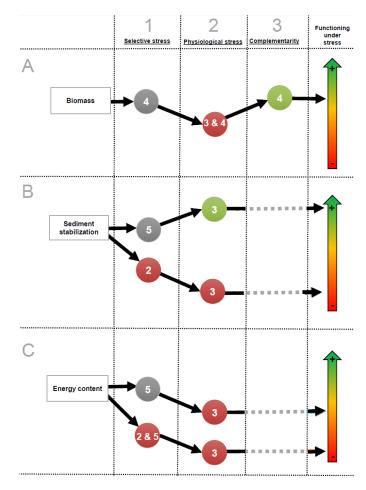
3236 The functional impact of selective biodiversity loss relative to changes in complementarity 3237 increases with differences in the species' functional contribution (Jiang 2007): when species 3238 contribute similarly to functioning, selective biodiversity loss can be compensated by an increase 3239 in complementarity effects, whilst this potential for compensation decreases when biodiversity 3240 loss selectively targets highly productive species. Complementarity commonly drives biomass 3241 production, but tends to have weaker effects on certain biochemical or trophic processes (Duffy 3242 2002, Jiang et al. 2008, Cardinale et al. 2011). Processes such as decomposition, trophic energy 3243 transfer, nutrient cycling, bioturbation or ocean carbon export can instead depend on the 3244 presence of organisms with a disproportionately high contribution to functioning (Emmerson et 3245 al. 2001, Hillebrand and Cardinale 2004, Waldbusser et al. 2004, Solan et al. 2004, Jiang 2007,

Jiang et al. 2008, Cardinale et al. 2011, Bracken and Low 2012, Van Colen et al. 2012, Litchman etal. 2015).

3248 Here, differences in functional contribution amongst diatom species were limited in terms of 3249 biomass production, and most pronounced in terms of energy content. The monoculture biomass 3250 of the most productive species (Nitzschia sp. in Chapter 2, N. arenaria in Chapters 4 and 5, 3251 respectively) exceeded that of the least productive species (S. robusta and A. lineolata) by a factor 3252 three and four, respectively. The species contributing most to sediment stabilization (*Nitzschia* sp. 3253 and A. lineolata in Chapters 2 and 5, respectively) produced seven times more EPS than the least 3254 productive species (S. robusta and N. arenaria). The most lipid-rich species (two different 3255 Nitzschia sp.) however produced fourty and fifteen times more fatty acids than the most lipid-poor 3256 species in the diatom communities of Chapters 2 and 5 (*N. arenaria* and *C. closterium*).

3257 In accordance with these differences in functional contribution, biomass production did not 3258 depend on the identity of the dominant species, and was reduced by physiological atrazine and 3259 copper effects, which could however be compensated by positive complementarity effects on 3260 stressed diatom communities (Fig. 6.2 A). The extent to which complementarity could compensate 3261 stress effects on diatom biomass production depended on the stress level, and whether 3262 complementarity favoured the growth of low- or high-yield species. For energy content and 3263 sediment stabilization, complementarity effects could not be quantified, since additive 3264 partitioning would require information on the fatty acid and EPS production of every species in 3265 monoculture and in community. Fig. 6.2 consequently does not make a statement on 3266 complementarity as driver of diatom energy content and sediment stabilization under stress.

3267 EPS was affected by selective stress, when stress-sensitive and tolerant species differed in their 3268 functional contribution. At low stress levels, diatoms increased their EPS production as a 3269 physiological stress response, whilst at high stress diatoms could not maintain their EPS 3270 production (Fig. 6.2 B). In a best-case scenario, stress thus did not target the functionally most 3271 important species, diatoms increased their EPS production at low stress, and stressed diatom communities produced more EPS than under control conditions. In a worst-case scenario, stress 3272 3273 targeted the most productive species and diatoms could not maintain their EPS production under 3274 high stress, causing a disproportionate loss of EPS production in stressed diatom communities 3275 (Fig. 6.2 B).



3278 Fig. 6.2: Summary of three effects (Selective stress - Panel 1, Physiological stress - Panel 2, 3279 Complementarity – Panel 3) determining the contribution of diatom communities to three ecosystem 3280 processes under stress (Biomass production – Panel A, Sediment stabilization – Panel B, Energy content – 3281 Panel C). Red, grey and green circles respectively represent negative, neutral and positive impacts of each effect on ecosystem processes. The numbers within the circles indicate the chapters in which the respective 3282 3283 effects were analysed (Chapters 2 to 5). The coloured arrows on the right-hand side of panels A, B and C represent an approximation to which extent stressed diatom communities could maintain their 3284 3285 contribution to ecosystem functioning (green: high functioning under stress; red: low functioning under 3286 stress). For the sake of simplicity this figure does not present the extent of ecosystem functioning change in 3287 the different treatments of Chapters 2 to 5. Instead, this figure has to be understood as a qualitative rather 3288 than quantitative interpretation of the impact of selective stress, physiological stress and complementarity 3289 on the contribution of stressed diatom communities to ecosystem functioning. Whilst the impact of selective 3290 stress, physiological stress and complementarity on diatom biomass production were consistently neutral, 3291 negative and positive respectively, changes in diatom energy content and sediment stabilization differed 3292 depending on stress type (atrazine or copper) and stress level. For diatom energy content and sediment 3293 stabilization, this figure thus indicates a best-case and worst-case scenario, resulting from the combination 3294 of selective and physiological atrazine and copper effects on the functional contribution of diatom 3295 communities.

3296

Diatom energy content was most affected by selective atrazine stress, due to the large differences in the species' fatty acid production (but see above). As predicted from the species' EC50 values, the most copper-tolerant species also produced most fatty acids, and selective copper stress had no effect on diatom energy content. Compared to biomass production and sediment stabilization, diatom energy content was also most affected by physiological stress, with the fatty acid 3302 production of most species being reduced at stress levels which did not affect other processes.
3303 Context-dependent effects of both stressors thus also caused a loss of diatom energy content, even
3304 in the absence of selective stress effects (Fig. 6.2 C). In a best-case scenario, diatom energy content
3305 was not affected by selective copper stress, and only reduced by context-dependent effects at high
3306 copper levels. In a worst-case scenario, low atrazine levels reduced diatom energy content
3307 through selective changes in community structure, whilst high levels of the herbicide further
3308 reduced diatom energy content through context-dependent effects.

3309 These strong effects of chemical stress on diatom energy content which, partly despite invariant 3310 biomass, knocked on to the next trophic level highlight the relevance of diet quality in a trophic 3311 BEF context. Trophic BEF research has mostly focussed on diet quantity, i.e. the biomass 3312 production of primary producers (Balvanera et al. 2006, Duffy et al. 2007). Diet quality received 3313 broad attention in marine trophic ecology when population declines of some well-known marine 3314 top predators could be explained by changes in the quality of their diet. For example, reproductive 3315 failures in seabirds (Wanless et al. 2005, Kitaysky et al. 2006, Österblom et al. 2008) and declines 3316 in sea lion populations (Anderson and Piatt 1999, Rosen and Trites 2005, Trites et al. 2007) have 3317 been linked to losses in the energy content of their diet, and to shifts in prey composition from 3318 lipid-rich to lipid-poor prey species.

3319 However, diet quality in terms of energy content does not only play a crucial role at the top of the 3320 food web, but is the key driver of energy transfer at the primary producer-consumer interface. In 3321 aquatic ecosystems, algal diet quality is highly sensitive to changing environmental conditions, 3322 and energy content shows considerably variation among and within algal classes (Dunstan et al. 1993, Taipale et al. 2013, Guo et al. 2016). Community shifts from high- to low-energy primary 3323 3324 consumers can cause trophic decoupling, i.e. low energy transfer efficiencies to the consumer level 3325 despite high producer biomass, which reduces consumer energy content and ultimately 3326 reproduction (Müller-Navarra et al. 2000, Guo et al. 2016). The limited duration of the 3327 experiments in this work allowed to test for the former (energy transfer) but not the latter 3328 (consumer reproduction) consequence of altered diet quality. Nevertheless, anthropogenic 3329 change is increasingly altering the energy content of primary producers (Teoh et al. 2013, Guo et 3330 al. 2016) and is causing large-scale shifts in both local and global phytoplankton compositions 3331 (e.g. from energy-rich diatoms to energy-poor cyanobacteria, Bopp et al. 2005, Moran et al. 2010, 3332 Litchman et al. 2015). Evaluating the functional impact of these changes not only in terms of diet 3333 quantity, but also diet quality and trophic energy transfer will be major task for future BEF 3334 research.

## 3335 **6.5. Future perspectives**

This doctoral thesis sets in many ways a baseline for the incorporation of anthropogenic stress into future research on biodiversity and ecosystem functioning research. The outcome of this work poses several challenges to further increase the level of realism in diversity-functioning research.

- 3340 A first challenge is to test which level of diversity is most relevant for ecosystem • 3341 functioning under stress. In this work, stress consistently altered evenness, which was a 3342 better predictor of functioning than richness (but see 6.1), whilst most BEF experiments 3343 to date have focused on the functional role of species richness (Wittebolle et al. 2009, 3344 Cardinale et al. 2011, Tilman et al. 2014). Describing diversity at the species level however 3345 only stands in the middle of a broad diversity spectrum, which includes genetic diversity 3346 (Hughes et al. 2008), functional trait diversity (Díaz et al. 2003, Cadotte et al. 2011) and 3347 the number of distinct communities within a system (Pasari et al. 2013).
- To quantify which type of diversity is most relevant for ecosystem functioning under stress, future BEF research would need experiments which explicitly manipulate biodiversity at multiple levels. For example, is ecosystem functioning under stress driven by genetic diversity (e.g. through the presence of tolerant genotypes), trait diversity (e.g. the number of effect traits and their correlation with response traits), species diversity or community diversity (e.g. the number of distinct communities in a stressed system)?
- A second challenge is to include multiple anthropogenic stressors into BEF experiments.
   The few BEF experiments which included explicit exposure to anthropogenic stressors
   examined the impact of one type of stress in isolation (McMahon et al. 2012, Steudel et al.
   2012, Radchuk et al. 2016, including this work). Ecosystems are however commonly
   exposed to multiple stressors (Lawler et al. 2006, Malaj et al. 2014).
- 3360 Future BEF research will need approaches to predict community structure and 3361 ecosystem functioning in the presence of multiple anthropogenic stressors. 3362 Community ecotoxicology has developed advanced mechanistic and predictive 3363 approaches to predict biodiversity in the presence of multiple contaminants 3364 (Koelmans et al. 2001, Rohr et al. 2006, Clements and Rohr 2009, Halstead et al. 3365 2014). Explicitly linking these multistressor approaches with the species' 3366 functional contribution represents a promising basis for testing the functional 3367 impact of multiple stressors, to further increase realism in BEF designs.

A third challenge for future BEF research is to examine not only the functional
 consequences of species loss, but also of species gain. BEF research has been motivated by

concerns about biodiversity loss, but global change, through species invasions and range
expansions, is homogenizing the distribution of organisms worldwide (Cardinale et al.
2012, Dornelas et al. 2014). While global biodiversity indeed seems to decrease,
biodiversity at local scales has been observed to remain unaffected or even increase (Sax
and Gaines 2003, Dornelas et al. 2014, McGill et al. 2015).

- In order to keep pace with a changing world, BEF research will need to develop designs that predict ecosystem functioning under simultaneous biodiversity losses and gains. The marine environment is largely affected by the homogenization of Earth's biota (Molnar et al. 2008, Vansteenbrugge et al. 2016), and coastal ecosystems are particularly vulnerable to invasive species (Carlton 1998, Gamfeldt et al. 2015). Marine studies could thus lead the way in integrating simultaneous invasion and extinction into BEF frameworks.
- 3382 Last, BEF research should further increase realism by including the spatial and temporal 3383 complexity driving the BEF relation in natural ecosystems. There still remains a large gap 3384 between the small spatial and temporal scales at which most BEF experiments are being 3385 performed, and those scales at which conservation efforts are taking place (Srivastava and 3386 Vellend 2005, Cardinale et al. 2011). Larger spatial and temporal scales allow for a greater 3387 niche partitioning, a key mechanism for complementarity (Cardinale et al. 2007, 3388 Stachowicz et al. 2008a, Venail et al. 2010). Facilitative interactions also grow stronger 3389 with time and can change with fluctuating abiotic conditions (Mulder et al. 2001, Cardinale 3390 et al. 2007, Stachowicz et al. 2008b).
- 3391 This discrepancy of experimental and natural systems highlights the need to cross-3392 check experimental results with observed patterns in natural ecosystems, and to 3393 transform the results of short-term laboratory or mesocosm BEF experiments into 3394 meaningful predictions at scales which matter to conservation. To extrapolate the 3395 results of controlled short-term experiments to more complex environments, BEF 3396 experiments should test biodiversity effects on functioning under constant 3397 conditions as well as different regimes of temporal and abiotic fluctuations. Once 3398 such an extrapolation degree has been established for various ecosystem 3399 processes, we should be able to transform the outcome of controlled small-scale 3400 BEF experiments to scales which are relevant for management and conservation 3401 efforts.

In a world affected by human activities at an increasing scale and speed, BEF research has come a
long but fast way, from being a 'niche research area' at the interface of community and ecosystem
ecology only twenty years ago to having a broad impact on both ecological research and public
awareness. BEF research went through a contentious confirmatory phase in the 1990s and an

3406 exploratory phase in the 2000s revealing the core drivers of the biodiversity effect in functioning.
3407 Now, BEF research faces a realism phase with the challenge to include the natural and
3408 anthropogenic drivers of diversity-functioning relations into its experimental design. By
3409 complementing current BEF designs with the inherent variability of natural systems and the
3410 anthropogenic drivers of the ongoing biodiversity change, we should be able to gain a more
3411 thorough understanding of ecosystem functioning.

## 3414 Addendum I – Supporting Material Chapter 2

Species	Shape	Biovolume equation	Length (a) [µm]	Width (b) [µm]	Height (c) [µm]	Biovolume [µm <sup>3</sup> ]
Navicula arenaria	Elliptic prism	$V = \frac{\pi}{4} \times a \times b \times c$	47	18	15	9896
Entomonei s paludosa	Elliptic prism	$V = \frac{\pi}{4} \times a \times b \times c$	40	5	50	7854
Seminavis robusta	cymbelloid	$V = \frac{1}{6}\pi \times (2b)^2 \times a \times \frac{\beta}{360};$	25	15	10	2813
Nitzschia sp.	Prism on parallelogram	with $sin \frac{\beta}{2} = \frac{c}{2 \times b}$ $V = \frac{1}{2} \times a \times b \times c$	45	15	17	5738

3418 Table S1: Cell biovolume per species. Biovolume was calculated based on the closest approximation of3419 geometric shape and the mean measured linear dimensions during the experiments.

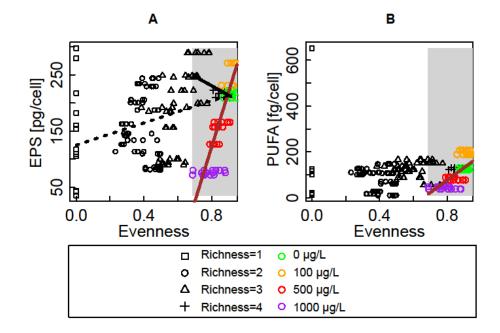


Fig. S1: EPS (A) and PUFA (B) concentrations normalized to cell density as a function of evenness.
Regression lines show significant relations between evenness and function. Brown lines are linear models
based on the atrazine experiment. Dashed and solid black lines are linear models based on the random
assembly experiment for the whole evenness range or for the evenness range of the atrazine experiment
(grey area), respectively.

	Model	Slope	s.e.	Т	Р	AIC	Log lik.
R	1	78,22	18,45	4,24	<0,0001	2029	-1012
	2	98,30	18,30	5,37	<0,0001	2023	-1008
А	1	971,96	98,94	9,82	<0,0001	781	-387
	2	894,93	83,61	10,70	<0,0001	780	-386
Rʻ	1	-188,23	81,60	-2,31	0,0309	234	-114
	R R n: A 1]	Data Model R 1 2 A 1 I] 2	Data         Model         Slope           R         1         78,22           2         98,30           4         1         971,96           1         2         894,93	Data         Model         Slope         s.e.           R         1         78,22         18,45           2         98,30         18,30           4         1         971,96         98,94           1         2         894,93         83,61	Data         Model         Slope         s.e.         T           R         1         78,22         18,45         4,24           2         98,30         18,30         5,37           A         1         971,96         98,94         9,82           1]         2         894,93         83,61         10,70	Data         Model         Slope         s.e.         T         P           R         1         78,22         18,45         4,24         <0,0001	Data         Model         Slope         s.e.         T         P         AIC           R         1         78,22         18,45         4,24         <0,0001

### 2 -166,25 64,94 -2,56 0,0179 221 -107 Yes 14,87 **0,0001** R 1 -25,99 31,09 -0,84 0,4043 2217 -1106 No 2 61,55 17,88 3,44 0,0007 2052 -1022 No NA NA Energy А 1 663,37 74,62 8,89 < 0,0001 740 -367 Yes content: 2 56,28 541,24 9,62 <0,0001 736 -364 0,0160 Yes 5,80 PUFA [fg/ml] Rʻ 1 -37,56 96,31 -0,39 0,7003 242 -118 No 2 63,13 63,25 1,00 0,3291 222 -107 No NA NA

Validity

Yes

Yes

Yes

Yes

Yes

LR

7,78

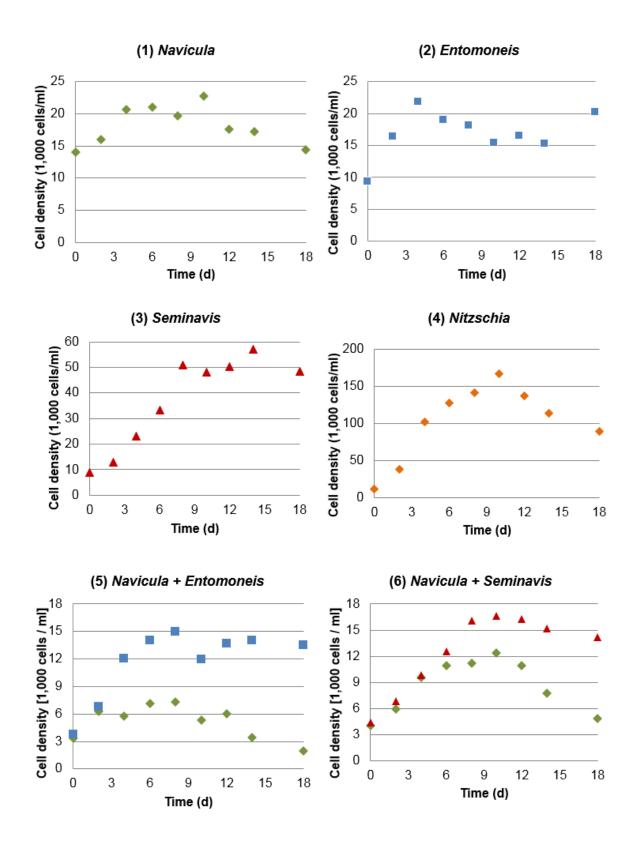
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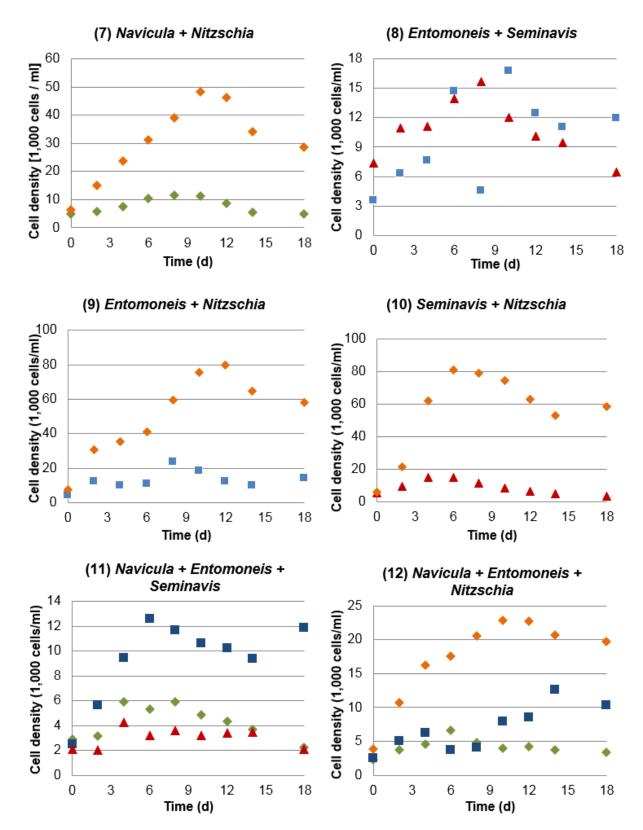
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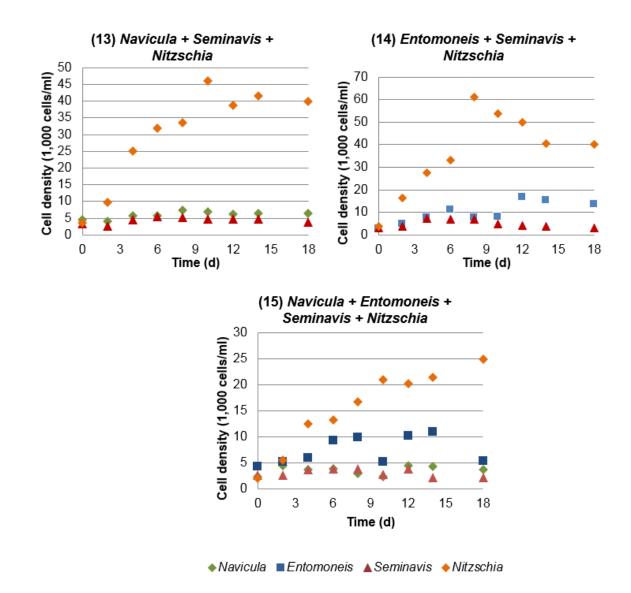
0,005

0,0897

3429 Table S2: Results of generalized least squared models predicting PUFA and EPS production (normalized to 3430 cell density) from evenness, without (model 1) or with (model 2) a variance structure. 'Data' denotes the 3431 data on which the models are based: the random assembly experiment (R), the atrazine experiment (A), the 3432 random assembly experiment for evenness values overlapping with those from the atrazine experiment 3433 (R'); 's.e.' is the standard error on the estimated slopes; T and P denote the t- and p-values, bold values are 3434 statistically significant. AIC is the Akaike information criterion; 'Validity' denotes if residuals were 3435 homogeneous and normally distributed ('yes') or not ('no'); 'LR' is the likelihood ratio of model 1 vs. model 3436 2, P LR the corresponding p-value.







**Fig. S2:** Cell density per species in the random assembly experiment. Treatments 1-15 (n=12) represent all

3441 possible random species combinations. Colour codes refer to the species *Navicula arenaria* (green), 3442 *Entomoneis paludosa* (blue), *Seminavis robusta* (red) and *Nitzschia sp.* (orange).

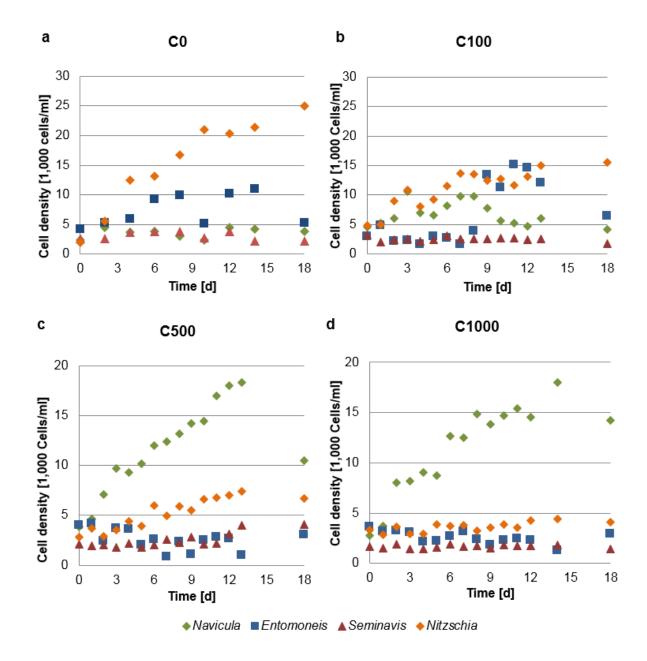


Fig. S3: Cell density per diatom species in the atrazine experiment. Treatments C0, C100, C500 and C1000
(0, 100, 500 and 1000 μg/l atrazine, n=18) are indicated in panels a, b, c and d respectively. Colour codes
refer to the species *Navicula arenaria* (green), *Entomoneis paludosa* (blue), *Seminavis robusta* (brown) and

*Nitzschia sp.* (orange).

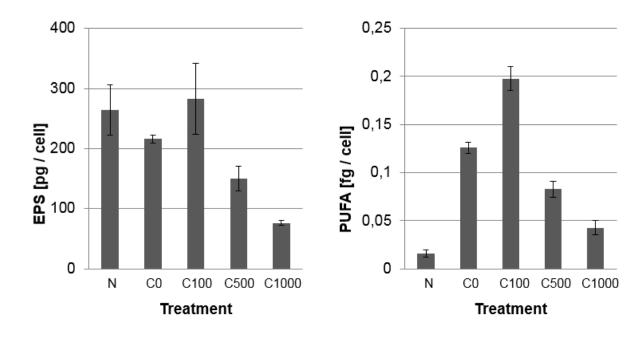




Fig. S4: EPS (a) and PUFA (b) concentrations normalized to cell density in monocultures of *Navicula arenaria* and atrazine-exposed communities. Treatments: N = *Navicula arenaria* monoculture, C0, C100, C500 and C1000 = full species community exposed to 0, 100, 500 and 1000 μg/l atrazine. Treatments C500 and C1000 were dominated by *Navicula arenaria* (s. Fig. S3). EPS: n=3, PUFA: n=2

# 3455 Addendum II – Supporting Material Chapter 3

Species	Acc. Nr.	K [10 <sup>3</sup> cells/ml]	Biovolume [µm3]	S:V	Length [µm]	Chl. a [F0]	TN [%]	ТОС [%]	Mix 0.1	Mix 0,6	PE [Fv/Fm]	EPS total [mg/ml]	EPS / cell [ng/cell]	FA total [μg/ml]	FA / cell [pg/cell]
Cylindrotheca	DCG	997	67	1.57	18.25	0.34	6.02	41.67	0.91	1.00	0.27	1.32	3.16	931.10	294.78
fusiformis	0423	(+- 32)	(+- 7)	(+- 0.08)	(+-1.0)	(+- 0.05)	(+- 1.2)	(+- 0.3)	(+- 0.1)	(+-0.1)	(+- 0.02)	(+- 0.01)	(+- 0.07)	(+- 310)	(+-103)
Nitzschia	DCG	170	380	0.61	26.15	0.27	NA	17.71	0.89	1.13	0.46	1.28	16.56	3.67	58.84
sp. 1	0483	(+- 26)	(+- 48)	(+- 0.04)	(+-2.1)	(+- 0.09)			(+- 0.2)	(+- 0.4)	(+- 0.02)	(+- 0.07)	(+-2.86)	(+-2.34)	(+-23)
Nitzschia	DCG	412	517	0.54	33.01	0.42	7.26	33.20	1.22	1.10	0.56	1.96	9.99	7.00	39.24
acicularis	0459	(+- 46)	(+- 68)	(+- 0.03)	(+-2.5)	(+- 0.07)	(+- 0.6)	(+- 2.5)	(+- 0.1)	(+-0.1)	(+- 0.02)	(+- 0.13)	(+- 0.32)	(+- 1.63)	(+-8)
Amphora	DCG	121	778	0.93	26.39	0.30	9.03	43.84	0.18	0.81	0.64	3.61	89.09	29.99	767.20
lineolata	0478	(+- 25)	(+- 123)	(+- 0.05)	(+- 1.5)	(+- 0.09)			(+- 0.1)	(+- 0.3)	(+- 0.04)	(+- 0.31)	(+- 0.72)	(+- 5.22)	(+-228)
Cylindrotheca		141	345	0.86	25.46	0.43	5.28	14.74	2.36	3.82	0.53	3.39	67.02	19.02	313.14
closterium		(+- 20)	(+- 60)	(+- 0.06)	(+- 1.0)	(+- 0.03)	(+- 1.7)	(+- 16)	(+- 0.3)	(+- 1.1)	(+- 0.04)	(+- 0.34)	(+- 4.48)	(+- 2.31)	(+-35)
Gyrosigma	DCG	42	2081	0.34	44.58	1.45	7.41	17.59	1.87	1.85	0.64	1.89	157.19	10.99	820.91
sp. 2	0491	(+- 6)	(+- 241)	(+- 0.02)	(+-2.2)	(+- 0.24)	(+-2.8)	(+- 3.4)	(+- 0.5)	(+- 0.4)	(+- 0.01)	(+- 0.21)	(+- 25.46)	(+- 3.62)	(+-207)
Navicula	DCG	58	4466	0.39	39.49	1.72	0.07	00.40	0.16	0.61	0.52	3.35	279.30	10.90	782.41
arenaria str. A2	0487	(+- 11)	(+- 388)	(+- 0.02)	(+- 2.8)	(+- 0.35)	2.86	22.19	(+- 0.1)	(+- 0.3)	(+- 0.04)	(+- 0.39)	(+- 15.80)	(+- 4.91)	(+- 155)
Astartiella	DCG	86	1552	0.59	33.19	2.71	7.49	31.90	0.89	0.92	0.63	2.88	84.10	21.34	687.92
bahusiensis	0469	(+- 28)	(+-239)	(+- 0.05)	(+- 0.3)	(+- 1.22)	(+-2.9)	(+- 8.9)	(+- 0.2)	(+- 0.2)	(+- 0.02)	(+- 0.51)	(+- 9.75)	(+- 2.59)	(+- 41)
Nitzschia	DCG	234	650	0.40	23.69	1.46	6.51	36.66	0.98	1.08	0.65	3.12	76.48	49.51	895.53
sp. 2	0421	(+- 22)	(+- 46)	(+- 0.02)	(+- 1.4)	(+- 0.20)	(+- 0.2)	(+- 2.9)	(+- 0.3)	(+- 0.3)	(+- 0.01)	(+- 0.69)	(+- 15.01)	(+- 5.13)	(+- 182)
Gyrosigma	DCG	120	2147	0.35	56.08	1.14	6.13	32.29	0.41	0.90	0.51	0.57	11.74	12.22	247.84
<i>sp.</i> 1	0468	(+- 12)	(+- 428)	(+- 0.04)	(+- 3.4)	(+- 0.19)	(+- 1.2)	(+- 1.4)	(+- 0.1)	(+- 0.3)	(+- 0.03)	(+- 0.24)	(+- 5.38)	(+- 4.79)	(+- 85)
Navicula	DCG	67	3120	0.46	39.80	1.62	8.69	25.19	0.78	0.38	0.45	2.39	80.98	8.95	282.64
<i>arenaria</i> str. A7	0489	(+- 10)	(+- 519)	(+- 0.04)	(+- 0.4)	(+- 0.37)	(+- 1.2)	(+- 6.4)	(+- 0.4)	(+- 0.2)	(+- 0.04)	(+- 0.13)	(+- 6.31)	(+- 1.41)	(+- 24)
Navicula	DCG	38	5738	0.34	35.50	2.39	13.90	17.71	1.40	1.05	0.49	3.91	254.63	7.93	508.04
digitoradiata	0490	(+- 6)	(+- 640)	(+- 0.02)	(+- 1.0)	(+- 0.72)	10.70	1/./1	(+- 0.5)	(+- 0.3)	(+- 0.01)	(+- 0.79)	(+- 51.01)	(+- 1.60)	(+- 46)

Navicula	DCG	258	587	0.79	21.46	0.65	8.10	31.14	0.79	0.79	0.37	2.16	17.31	14.89	144.67
sp. 2	0671	(+- 34)	(+- 113)	(+- 0.05)	(+- 1.6)	(+- 0.12)	(+- 0.1)	(+-1.4)	(+- 0.3)	(+- 0.2)	(+- 0.03)	(+- 0.46)	(+- 3.41)	(+- 9.04)	(+-87)
Amphora	DCG	69	533	1.01	24.21	1.58	11.45	21.25	0.75	0.42	0.58	3.56	158.77	5.52	215.01
sp. 1	0481	(+- 16)	(+- 87)	(+- 0.06)	(+- 1.3)	(+- 0.38)	(+- 2.9)	(+- 15)	(+- 0.2)	(+- 0.2)	(+- 0.05)	(+- 0.27)	(+- 29.13)	(+- 0.31)	(+- 58)
Biremis	DCG	15	4108	0.37	29.39	2.99	14.21	28.18	1.25	1.50	0.43	1.51	203.54	2.24	377.72
ambigua	0480	(+-3)	(+- 593)	(+- 0.02)	(+- 1.5)	(+- 0.66)	14.21	20.10	(+- 0.4)	(+- 0.7)	(+- 0.05)	(+- 0.12)	(+- 29.55)	(+- 0.41)	(+- 57)
Entomoneis	DCG	45	10932	0.28	41.53	2.97	17.52	22.79	1.01	0.96	0.60	0.59	39.20	20.78	1292.26
sp.	0466	(+- 12)	(+-2767)	(+- 0.03)	(+- 3.3)	(+-0.71)	(+-10)	(+- 9.9)	(+- 0.3)	(+- 0.2)	(+- 0.02)	(+- 0.03)	(+- 8.05)	(+- 3.27)	(+- 489)
Navicula		302	303	1.02	19.06	1.11	7.37	26.21	1.40	1.54	0.51	2.19	16.29	8.86	67.91
sp. 1		(+- 46)	(+- 80)	(+- 0.12)	(+- 0.5)	(+- 0.24)	(+- 1.5)	(+- 1.9)	(+- 1.0)	(+- 0.6)	(+- 0.02)	(+- 0.49)	(+- 4.47)	(+- 7.91)	(+- 60)
Amphora	DCG	132	1052	0.88	24.57	2.22	7.97	28.24	1.09	0.78	0.56	0.67	11.03	5.69	78.96
sp. 2	0672	(+-7)	(+- 183)	(+- 0.10)	(+- 1.2)	(+- 0.12)	(+- 0.5)	(+-11)	(+- 0.2)	(+- 0.2)	(+- 0.03)	(+- 0.2)	(+- 3.35)	(+- 0.42)	(+-7)

3459

3460 **Table S1:** Listing of the 18 diatom strains used in the experiments. From left to right are indicted: species name, strain accession number (Acc. Nr.) in the diatom

3461 culture collection (BCCM/DCG) of the Belgian Coordinated Collection of Micro-organisms (<u>http://bccm.belspo.be</u>), carrying capacity K (n=6), traits and functional

3462 contribution as determined in the control treatments of the atrazine experiment: biovolume (n=12), surface to volume ratio (S:V, n=12), length (n=12), chlorophyll a

3463 content (Chl. a, n=6; minimum chlorophyll fluorescence signal F<sub>0</sub> per 10<sup>6</sup> cells), % total nitrogen (TN, n=2 or 1), % total organic carbon (TOC, n=2 or 1), mixotrophy 3464 (Mix0.1 and Mix0.6, n=6, change of carrying capacity in the presence of 0.1 mM and 0.6 mM Glucose), photosynthetic efficiency (PE, n=6, ratio of variable to maximum

3465 fluorescence), EPS (n=3) per ml culture and per cell, polyunsaturated fatty acids (FA, n=2) per ml culture and per cell.

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Substance	Theoretical concentration [µg/l]	Measured concentration [µg/l]
Control	0	<1
Atrazine	50	48
Atrazine	100	91
Atrazine	200	190
Atrazine	500	542
Atrazine	1000	1080
Copper	30	29
Copper	90	94
Copper	270	271
Copper	810	826

**Table S2:** Experimental atrazine and copper treatments with indication of theoretical and verified atrazine and copper concentrations as determined by GC-MS and graphite furnace AAS respectively (Limits of quantification : < 1,0  $\mu$ g/L atrazine and 2.6  $\mu$ g/L copper for 10 ml samples). 

Response	Data	Mod	Predictors	Slope	s.e.	Т	Р	AIC	Log lik.	Validity
Numerical:	Con	•	Biovolume	0.03	0.003	8.82	< 0.0001	1075	(24	V
Carrying	Cu	A	Mix	-31.6	23.4	-1.35	0.20	1275	-634	Yes
capacity	Atr	A	Mix	144.0	30.0	4.76	0.0003	190	-91.9	Yes
	Cu	•	Biovolume	0.12	0.01	6.73	<0.0001	1502	-787	Vea
Functional:	Cu	A	Mix	-110.9	115.8	-0.96	0.35	1582		Yes
Functional.		Δ	Biovolume	0.18	0.01	13.87	<0.0001	1400 726		N
Photosynthetic	A +	A	Mix	188.7	102.2	1.85	0.08	1480	-736	No
efficiency	Atr	Ъ	Biovolume	0.11	0.02	5.76	<0.0001	1200	COF	V
		В	Mix	187.7	51.8	3.62	0.0028	1399	-695	Yes
Functional:	Cu	Δ	Biovolume	0.056	0.007	6.75	<0.0001	607	-300	Yes
	Cu	A	Mix	-57.2	42.1	-1.36	0.20	607		
EPS	Atr	A	Mix	-46.7	31.9	-1.47	0.16	192	-92.8	Yes
			Biovolume	0.043	0.006	6.62	<0.0001	252	1 17 1	N
	Con	A	Mix	-11.1	19.8	-0.56	0.58	352	-171	No
Functional:	Cu	Ъ	Biovolume	0.051	0.011	4.47	0.0001	244	107	V
Fatty acids		В	Mix	-17.4	17.0	-1.02	0.33	344	-167	Yes
ratty actus	A +	Δ	Biovolume	0.028	0.008	3.52	0.0014	410	205	
	Atr	A	Mix	-26.1	34.6	-0.76	0.46	418	-205	Yes

3475

3476 Table S3: Results of generalized least squares models predicting diatom numerical and functional response 3477 to copper and atrazine from traits; 'Response' denotes the response variable: numerical or functional 3478 response in terms of photosynthetic efficiency, EPS or fatty acid production; 'Data' denotes the data on 3479 which the models are based: the copper experiment (Cu) or the atrazine experiment (Atr); 'Mod' indicates 3480 if the model was fitted without (Model A) or with (Model B) variance structure; Predictors are biovolume 3481 and mixotrophic response to 0.6 mM Glucose concentrations (Mix); 'Slope' indicates the relation between 3482 predictors and numerical and functional response to copper and atrazine; 's.e.' is the standard error on the 3483 estimated slopes; 'T' and 'P' denote the t- and p-values, bold values are statistically significant, italic values 3484 are statistically significant over the whole dataset but not if outlier species are omitted (model fits without outlier species are then indicated in Table S4); 'AIC' is the Akaike information criterion, 'Log lik' the log-3485 3486 likelihood ratio. 'Validity' denotes if residuals were homogeneous and normally distributed ('yes') or not 3487 ('no'). If 'no', models were refitted ('Model B') with a variance structure allowing the residuals to change 3488 with the predictor (the predictor along which residuals were allowed to vary is marked in italic). Only bestfitting models with uncorrelated predictors are shown (p<0.05, correlation factor < 0.5, sequential 3489 3490 removing of predictors, evaluation of model fit by AIC and log likelihood); Additional model fits with traits 3491 that also predicted numerical and functional response to copper and atrazine, but were correlated to 3492 biovolume and mixotrophy (correlation factor > 0.5), are indicated in Table S4.

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Response	Data	Mod	Predictors	Slope	s.e.	Т	Р	AIC	Log lik.	Validity
	<b>0</b> (	D	Biovolume	0.04	0.006	7.08	<0.0001	1100	505	
	Cuʻ	В	Mix	-28.0	26.8	-1.04	0.31	1199	-595	Yes
			SV	-206	$0$ $26.8$ $-1.04$ $0.31$ $11^{10}$ $6$ $37$ $-5.53$ $<0.0011$ $4$ $11.01$ $0.49$ $0.62$ $12^{10}$ $4$ $11.01$ $0.49$ $0.62$ $12^{10}$ $0.0011$ $12^{10}$ $6$ $30.2$ $-1.34$ $0.200$ $12^{10}$ $0$ $0.67$ $10.21$ $<0.0001$ $12^{10}$ $1$ $7.95$ $2.65$ $0.0093$ $12^{10}$ $3$ $9.80$ $-2.89$ $0.0119$ $17^{10}$ $7$ $64.1$ $2.52$ $0.0244$ $17^{10}$ $9$ $47.4$ $4.89$ $0.0002$ $17^{10}$ $0.5$ $0.009$ $0.54$ $0.59$ $12^{10}$ $8$ $145.1$ $-4.61$ $<0.0011$ $15^{10}$ $0$ $44.7$ $-1.99$ $0.049$ $15^{10}$ $9$ $117.3$ $-1.19$ $0.25^{10}$ $15^{10}$ $1$ $11.3$ $-1.18$ $0.26^{10}$ $15^{10}$ $1$ $11.3$ $-1.18$ $0.26^{10}$ $12^{10}$ $1$ $11.3$ $-1.18$ $0.0011$ $14^{10}$ $2$ $116.9$ $-1.81$ $0.0011$ $12^{10}$ $2$ $116.9$ $-1.81$ $0.0011$ $12^{10}$ $1$ $0.12^{10}$ $5.8^{10}$ $0.0011$ $14^{10}$ $2$ $116.9$ $-1.81$ $0.0001$ $14^{10}$ $1$ $0.2^{10}$ $0.0001$ $12^{10}$ $14^{10}$ $1$ $11.9^{10}$ $3.58$ $0.0017$ $14^{10}$ <t< td=""><td></td><td></td><td></td></t<>					
		В	Chl. a	5.44	11.01	0.49	0.62	1272	-630	Yes
Numerical:	0		Mix	-40.6	30.2	-1.34	0.20			
Carrying	Cu		Length	6.80	0.67	10.21	<0.0001			
capacity		В	Chl. a	21.1	7.95	2.65	0.0093	1236	-612	Yes
			Mix	-28.3	9.80	-2.89	0.0119			
	A. (	A	Mix	161.7	64.1	2.52	0.0244	177	-85.6	No
	Atr'	В	Mix	281.9	47.4	4.89	0.0002	178	-85.2	Yes
	Atr"	A	Biovolume	0.005	0.009	0.54	0.59	1260	-627	Yes
			SV	-668	145.1	-4.61	<0.0001			
		В	Chl. a	-89.0	44.7	-1.99	0.049	1589	-788	Yes
	0		Mix	-139	117.3	-1.19	0.25			
	Cu		Length	21.2	3.71	5.71	<0.0001			
		В	Chl. a	-49.5	41.1	-1.20	0.23	1587	-787	Yes
			Mix	-131	111.3	-1.18	0.26			
Functional:	<i>(</i>		Biovolume	0.11	0.02	5.26	<0.0001	1.100		
Photosynthetic – efficiency	Cu'	В	Mix	-212	116.9	-1.81	0.09	1486	-738	Yes
	• • "		Biovolume	0.09	0.02	6.37	<0.0001	4005	(20)	
	Atr"	В	Mix	185.7	51.9	3.58	0.0034	1287	-639	Yes
			SV	-336	42.7	7.99	< 0.0001	0.0001	202	
		В	Mix	129.4	59.7	2.17	9 <0.0001 1465	1465	-727	Yes
	Atr		Length	14.7	4.19	3.51	0.0007			
		В	Chl. a	81.2	25.1	3.24	0.0017	1433	-710	Yes
			Mix	180.7	61.7	2.93	0.0017			
			SV	-390	97.1	4.02	0.0002			
		A	Chl. a	-14.5	32.6	-0.44	0.66	607	-300	Yes
	0		Mix	-71.4	60.8	-1.17	0.26			
Functional:	Cu		Length	8.13	2.14	3.50	0.0011			
EPS		A	Chl. a	25.4	31.2	0.81	0.42	608	-299	Yes
210			Mix	-54.3	59.5	-0.91	0.388			
	<i>c</i> , <i>i</i>		Biovolume	0.058	0.013	4.50	0.0001	F00	207	
	Cu'	A	Mix	-56.6	45.2	-1.25	0.23	580	-287	Yes
			SV	-336	93.5	-3.60	0.0013			
		A	Chl. a	50.0	19.9	2.51	0.0185	336	-163	Yes
Functional:	_		Mix	-2.42	32.3					
Fatty acids	Cu		Length	4.02	1.41	2.85	0.0083			Yes
i arry actus		В	Chl. a	28.0	22.0	1.27	0.22	336	-162	
			Mix	-27.5	21.1	-1.30	0.22			

Cut	В	Biovolume	0.046	0.011	4.17	0.0013	316	-153	,
Cuʻ	D	Mix	-20.9	14.7	-1.43	0.18	510	-155	Y
Atr"	В	Biovolume	0.005	0.011	0.46	0.65	385	-188	Yes
Au	В	Mix0.6	-33.5	31.6	-1.06	0.31	303	-100	Ĭ

3497 Table S4: Results of generalized least squares models predicting numerical and functional response to 3498 copper and atrazine from traits; 'Response' denotes the response variable: numerical and functional 3499 response in terms of photosynthetic efficiency, EPS or fatty acid production; 'Data' denotes the data on 3500 which the models are based: the copper experiment (Cu) or the atrazine experiment (Atr); the copper 3501 experiment without the largest species (Entomoneis sp., (Cu')), the atrazine experiment without the most 3502 mixotrophic species (Cylindrotheca closterium, (Atr')), and the atrazine experiment without the largest 3503 species (Entomoneis sp., (Atr")); 'Mod' indicates if the model was fitted without (Model A) or with (Model 3504 B) variance structure; Predictors are biovolume, surface to volume ratio (SV), chlorophyll a content (Chl. a), 3505 Length and mixotrophic response to 0.6 mM Glucose concentrations (Mix); 'Slope' indicates the relation 3506 between predictors and numerical and functional response to copper and atrazine; 's.e.' is the standard 3507 error on the estimated slopes; Only models with uncorrelated predictors that predicted numerical and 3508 functional response are shown (p<0.05, correlation factor < 0.5, sequential adding and removing of 3509 predictors, evaluation of model fit by AIC and log likelihood ratio); 'T' and 'P' denote the t- and p-values, 3510 bold values are statistically significant, italic values are statistically significant over the whole dataset but 3511 not if outlier species are omitted; 'AIC' is the Akaike information criterion, 'Log lik' the log-likelihood. 3512 'Validity' denotes if residuals were homogeneous and normally distributed ('yes') or not ('no'). If 'no', 3513 models were refitted ('Model B') with a variance structure allowing the residuals to change with the 3514 predictor (the predictor along which residuals were allowed to vary is marked in italic).

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Species		EC50 [	µg/l]	
	Carrying capacity	Photosynthetic efficiency	EPS / cell	Fatty acids / cell
Cylindrotheca	217.19 (3.15)	221.29 (13.46)	220.05 (101.35)	23.62 (21.22)
fusiformis	54.94 (10.01)	75.77 (27.98)	40.91 (8.03)	
Nitzschia sp. 1	137.09 (40.09)	231.41 (42.18)	478.08 (229.89)	95.71 (3.90)
	96.46 (22.94)	712.15 (65.91)	157.14 (56.86)	<i>84.03 (7.80)</i>
Nitzschia acicularis	213.31 (33.58)	292.72 (23.14)	112.91 (10.87)	236.54 (11.01)
	281.12 (44.41)	1511.18 (299.76)	531.48 (172.67)	117.07 (79.21)
Amphora lineolata	175.08 (18.32)	391.50 (44.71)	192.31 (157.25)	66.54 (2.04)
	287.76 (131.17)	703.83 (98.68)	178.38 (47.59)	107.58 (18.31)
Cylindrotheca	652.28 (171.90)	987.78 (65.51)	59.59 (7.64)	4.78 (3.33)
closterium	39.14 (10.50)	256.40 (54.26)	31.19 (0.94)	13.71 (7.75)
<i>Gyrosigma</i> sp. 2	453.41 (159.01) 271.42 (98.76)	499.60 (32.72) 730.73 (15.75)	216.74 (88.37)	85.50 (79.62) 325.09 (115.22)
<i>Navicula arenaria</i>	349.01 (128.63)	1144.84 (257.17)	176.89 (16.70)	209.07 (57.34)
str. A2	219.22 (50.81)	1005.53 (325.52)	581.77 (41.40)	179.27 (1.95)
Astartiella	328.79 (84.83)	296.67 (26.56)	172.73 (27.45)	325.68 (18.47)
bahusiensis	243.28 (54.76)	479.63 (118.04)	158.30 (35.39)	163.35 (137.67)
Nitzschia sp. 2	161.77 (49.61)	292.78 (34.05)	76.94 (57.10)	47.63 (29.93)
	64.78 (17.83)	434.83 (78.93)	237.49 (54.50)	89.58 (32.25)
<i>Gyrosigma</i> sp. 1	337.01 (72.29)	759.70 (41.81)	253.62 (154.76)	134.51 (26.60)
	180.63 (58.23)	403.43 (22.28)	175.52 (42.58)	95.28 (3.14)
Navicula arenaria	171.36 (32.67)	275.68 (42.65)	379.90 (54.36)	68.40 (5.79)
str. A7	348.17 (152.21)	1267.94 (370.70)	515.57 (197.42)	288.86 (4.92)
Navicula	217.51 (120.77)	616.91 (77.63)	184.35 (64.45)	138.98 (57.13)
digitoradiata	258.05 (87.62)	955.84 (102.73)	380.21 (230.53)	
Navicula sp. 2	57.61 (16.99)	446.60 (89.33)	189.20 (12.23)	52.52 (66.30)
	70.01 (26.09)	157.66 (18.38)	258.57 (104.69)	189.24 (1.36)
Amphora sp. 1	230.13 (124.85)	551.16 (182.05)	113.50 (42.56)	205.04 (0.12)
	132.97 (46.56)	698.06 (29.55)	190.75 (85.02)	80.79 (91.09)
Biremis ambigua	415.03 (76.71) 134.41 (32.92)	581.31 (96.54) 144.12 (27.81)	178.98 (85.62) 205.49 (85.05)	119.23 (22.55)
Entomoneis sp.	329.94 (230.12)	2676.35 (746.98)	251.21 (98.69)	507.03 (185.09
	396.63 (61.51)	1934.08 (488.57)	792.79 (75.41)	502.39 (243.05
Navicula sp. 1	400.87 (101.04)	404.32 (48.76)	155.33 (28.56)	262.52 (231.74
	35.12 (12.05)	175.19 (13.18)	140.05 (33.51)	<i>3.36 (0.75)</i>
Amphora sp. 2	148.73 (15.07)	626.61 (33.71)		100.27 (9.66)
Average	288.32 (143.79)	627.70 (589.20)	200.71 (103.06)	150.72 (129.87
	181.27 (111.43)	631.83 (504.96)	299.87 (212.88)	141.89 (124.90)

Table S5: Atrazine and copper EC50s obtained for carrying capacity (n=6), photosynthetic efficiency (n=6),
 fatty acid (n=2) and EPS production (n=3) of 18 benthic diatom strains. The first line refers to the results
 with atrazine, the second with copper (italics). Standard deviations are indicated in brackets.

Function	Data	Mod	Predictors	Slope	s.e.	Т	Р	AIC	Log lik.	Validity
		A	EC50 D	2.7e-4	7.8e-5	3.44	0.0008	-166	86	Yes
_	Cu	Α	EC50 PE	6.7e-5	1.8e-5	3.69	0.0004	-165	86	No
Photo- synthetic		В	EC50 PE	4.0e-5	1.1e-5	3.52	0.0006	-178	93	Yes
efficiency	A 4		EC50D	4.8e-5	6.3e-5	-0.76	0.45	100	65	V
Atr	Atr	A	EC50 PE	3.3e-5	1.8e-5	1.84	0.07	-123	65	Yes
	C	A	EC50 D	0.091	0.026	3.48	0.0010	489	-241	Yes
FDC	Cu	А	EC50 EPS	0.039	0.014	2.71	0.0093	439	-217	Yes
EPS	A 4		EC50 D	0.011	0.078	0.14	0.89	(0)	200	V
	Atr	A	EC50 EPS	-0.121	0.106	-1.14	0.26	606	-299	Yes
	6	А	EC50 D	-0.043	0.220	-0.19	0.84	392	-193	Yes
Fatty	Cu	Α	EC50 FA	-0.011	0.187	-0.06	0.95	380	-187	Yes
acids	A 4-14	tr A	EC50 D	0.515	0.383	1.34	0.19	402		Vaa
	Atr		EC50 FA	0.764	0.457	1.67	0.11	492	-242	Yes

3523 Table S6: Results of generalized least squared models relating functional contribution to numerical and 3524 functional response to copper and atrazine. 'Function' denotes the response variable: photosynthetic 3525 efficiency and fatty acid and EPS production per cell in unstressed conditions; 'Data' denotes the data on 3526 which the models are based: the copper experiment (Cu) or the atrazine experiment (Atr); 'Mod' indicates 3527 if the model was fitted without (Model A) or with (Model B) variance structure; Predictors are numerical 3528 response (EC50 D) and functional response to copper and atrazine (EC50 PE for photosynthetic efficiency, 3529 EC50 EPS for EPS, EC50 FA for fatty acids); if uncorrelated, numerical and functional response were 3530 included as predictors in a multiple regression, if numerical and functional response were correlated, 3531 separate regressions were fitted; 'Slope' indicates the relation between numerical and functional response 3532 to copper and atrazine and functioning under control conditions; 's.e.' is the standard error on the estimated slopes; 'T' and 'P' denote the t- and p-values, bold values are statistically significant; 'AIC' is the Akaike 3533 information criterion, 'Log lik' the log-likelihood. 'Validity' denotes if residuals were homogeneous and 3534 3535 normally distributed ('yes') or not ('no'). If 'no', models were refitted ('Model B') with a variance structure 3536 allowing the residuals to change with the predictor (the predictor along which residuals were allowed to 3537 vary is marked in italic).

3538

FR	Data	Mod	Predictor	Slope	s.e.	Т	Р	AIC	Log lik.	Validity
Primary Production: Photosynth. efficiency	Cu	Α	EC50 K	3.11	0.29	10.75	<0.0001	1575	-784	Yes
		В	EC50 K	3.20	0.30	10.51	<0.0001	1535	-763	Yes
	Atr	А	EC50 K	1.03	0.34	3.05	0.0029	1582	-788	Yes
Sediment stabilization: EPS	Cu	А	EC50 K	1.26	0.23	5.50	<0.0001	630	-312	Yes
	Atr	А	EC50 K	-0.27	0.10	-2.74	0.0085	632	-313	Yes
		В	EC50 K	-0.21	0.04	-5.44	<0.0001	620	-306	Yes
Energy content: Fatty acids	Cu	А	EC50 K	0.78	0.18	4.40	0.0001	365	-180	No
		В	EC50 K	0.67	0.19	3.52	0.0015	360	-176	Yes
	Atr	А	EC50 K	-0.11	0.15	-0.78	0.44	431	-212	No
		В	EC50 K	-0.26	0.11	-2.32	0.0266	429	-211	Yes

3542 Table S7: Results of generalized least squared models relating numerical and functional response. 'FR' 3543 denotes the response variable: functional response to copper or atrazine in terms of photosynthetic 3544 efficiency, fatty acid or EPS production; 'Data' denotes the data on which the models are based: the copper 3545 experiment (Cu) or the atrazine experiment (Atr); 'Mod' indicates if the model was fitted without (Model A) 3546 or with (Model B) variance structure; Predictors are numerical response (the EC50 of carrying capacity K) to copper and atrazine respectively; 'Slope' indicates the relation between numerical and functional 3547 3548 response; 's.e.' is the standard error on the estimated slopes; 'T' and 'P' denote the t- and p-values, bold 3549 values are statistically significant; 'AIC' is the Akaike information criterion, 'Log lik' the log-likelihood. 3550 'Validity' denotes if residuals were homogeneous and normally distributed ('yes') or not ('no'). If 'no', 3551 models were refitted ('Model B') with a variance structure allowing the residuals to change with the 3552 predictor (the predictor along which residuals were allowed to vary is marked in italic).

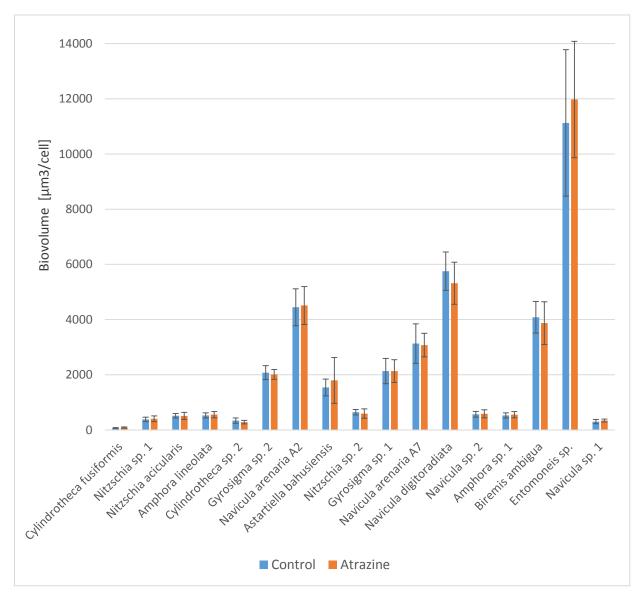
Species	Biovolume Control		Biovolun	ne Atrazine	Т	Р
Cylindrotheca fusiformis	68	(+-20)	84	(+-23)	-1.87	0.08
Nitzschia sp. 1	380	(+- 81)	406	(+- 102)	-0.14	0.88
Nitzschia acicularis	514	(+- 80)	508	(+- 127)	0.13	0.90
Amphora lineolata	527	(+- 91)	553	(+- 113)	-0.64	0.53
Cylindrotheca sp. 2	340	(+- 94)	281	(+- 64)	1.78	0.09
Gyrosigma sp. 2	2078	(+- 252)	2013	(+- 179)	0.73	0.47
Navicula arenaria A2	4447	(+- 670)	4510	(+- 687)	-0.23	0.82
Astartiella bahusiensis	1540	(+- 303)	1794	(+- 832)	-0.99	0.34
Nitzschia sp. 2	643	(+- 99)	593	(+- 171)	0.88	0.39
Gyrosigma sp. 1	2134	(+- 458)	2134	(+- 411)	< 0.01	0.99
Navicula arenaria A7	3131	(+-717)	3074	(+- 427)	0.23	0.82
Navicula digitoradiata	5753	(+- 694)	5316	(+- 762)	1.47	0.16
Navicula sp. 2	568	(+- 102)	585	(+- 145)	-0.33	0.75
Amphora sp. 1	527	(+- 91)	553	(+- 113)	-0.64	0.53
Biremis ambigua	4082	(+- 575)	3872	(+- 773)	0.76	0.46
Entomoneis sp.	11124	(+- 2654)	11976	(+- 2108)	-0.87	0.39
Navicula sp. 1	304	(+- 80)	342	(+- 50)	-1.41	0.18

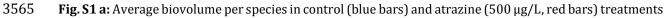
**Table S8 a:** Biovolume per species at the end of the experiment, in control and atrazine (500  $\mu$ g/L)3555treatments. T and P denote the t- and p-values from a two-sided t-test testing differences in biovolume in3556control vs. atrazine treatments, bold values are statistically significant (p<0.05).</td>

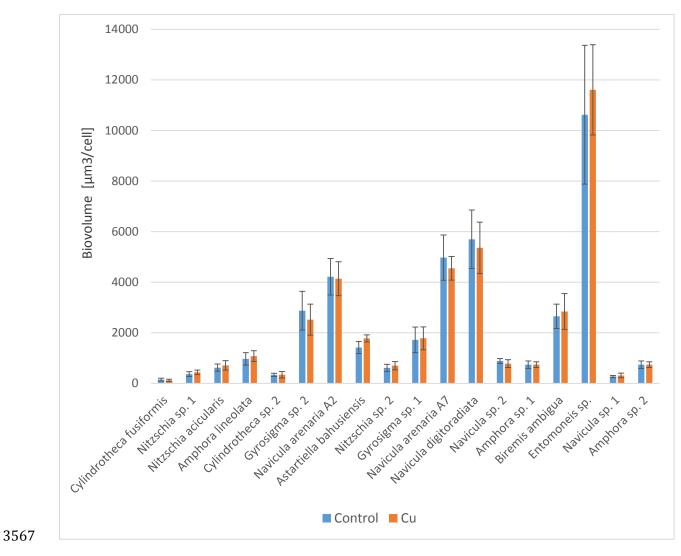
Species	Biovolume Control		Biovolume Cu		Т	Р
Cylindrotheca fusiformis	149	(+- 57)	122	(+- 44)	-1.30	0.21
Nitzschia sp. 1	366	(+- 95)	438	(+- 88)	-1.79	0.09
Nitzschia acicularis	622	(+- 144)	712	(+- 189)	-1.31	0.20
Amphora lineolata	967	(+- 247)	1080	(+- 213)	-1.20	0.25
Cylindrotheca sp. 2	334	(+- 64)	342	(+- 127)	-0.20	0.84
Gyrosigma sp. 2	2873	(+- 767)	2522	(+- 613)	1.24	0.23
Navicula arenaria A2	4214	(+- 723)	4139	(+- 672)	0.26	0.79
Astartiella bahusiensis	1418	(+- 242)	1779	(+- 140)	-4.48	0.0003
Nitzschia sp. 2	612	(+- 141)	702	(+- 160)	-1.45	0.16
Gyrosigma sp. 1	1722	(+- 509)	1782	(+- 454)	-0.30	0.76
Navicula arenaria A7	5077	(+- 1204)	4146	(+- 828)	1.45	0.17
Navicula digitoradiata	5698	(+- 1155)	5359	(+- 1015)	0.76	0.18
Navicula sp. 2	884	(+- 96)	780	(+- 154)	1.98	0.06
Amphora sp. 1	739	(+- 147)	740	(+- 114)	-0.02	0.98
Biremis ambigua	2650	(+- 482)	2843	(+- 707)	-0.78	0.45
Entomoneis sp.	10621	(+- 2743)	11607	(+- 1786)	-1.04	0.31
Navicula sp. 1	270	(+- 43)	313	(+- 93)	-1.45	0.17
Amphora sp. 2	739	(+- 147)	740	(+- 114)	-0.02	0.98

**Table S8 b:** Biovolume per species at the end of the experiment, in control and copper (270  $\mu$ g/L) treatments. T and P denote the t- and p-values from a two-sided t-test testing differences in biovolume in control vs. atrazine treatments, bold values are statistically significant (p<0.05).

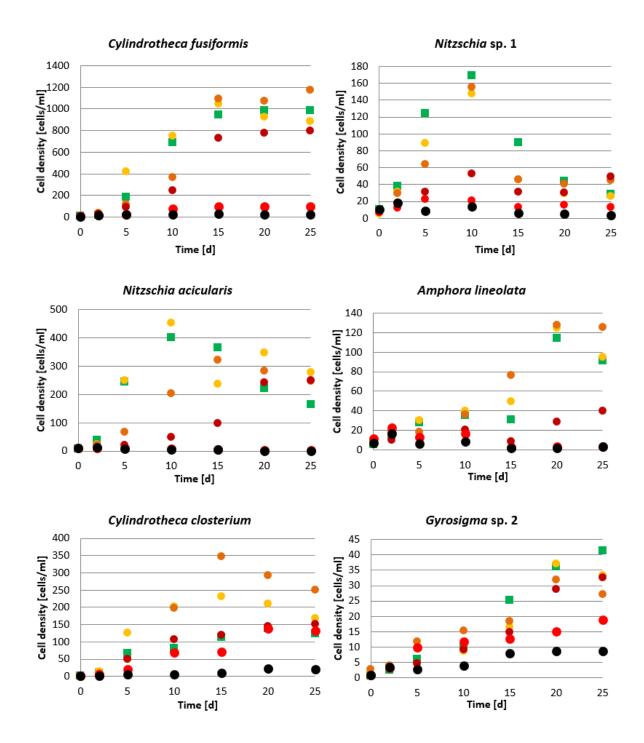


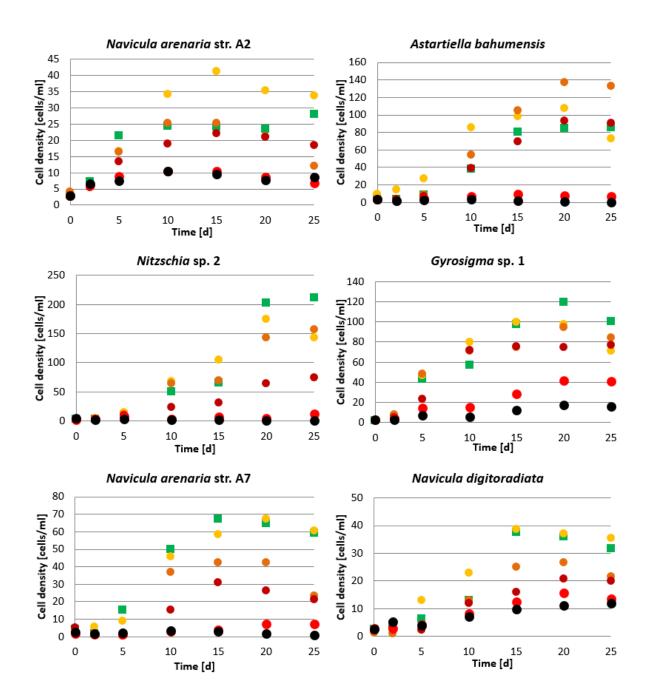




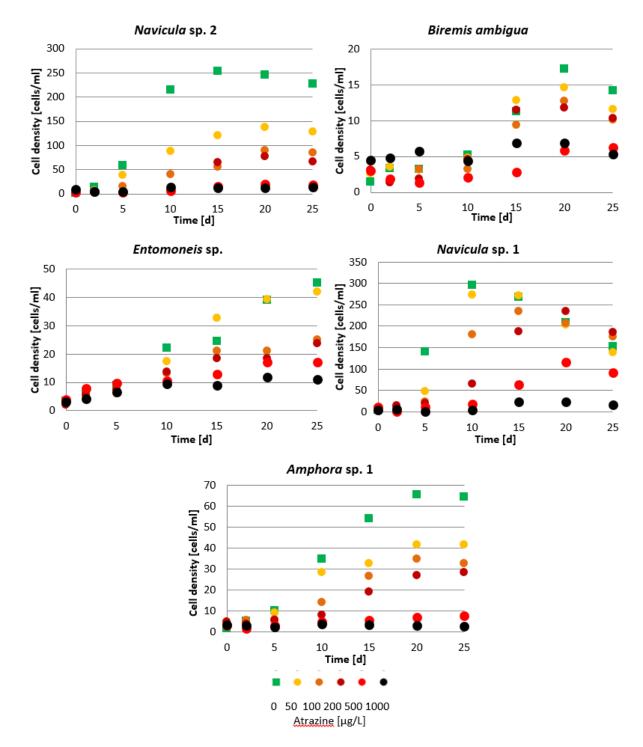


**Fig. S1 b.** Average biovolume per species in control (blue bars) and copper (270 μg/L, red bars) treatments

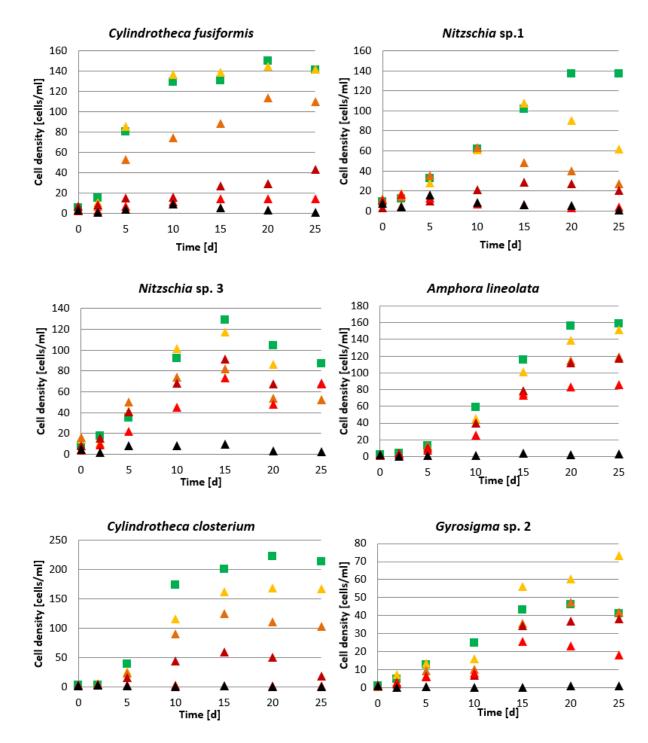


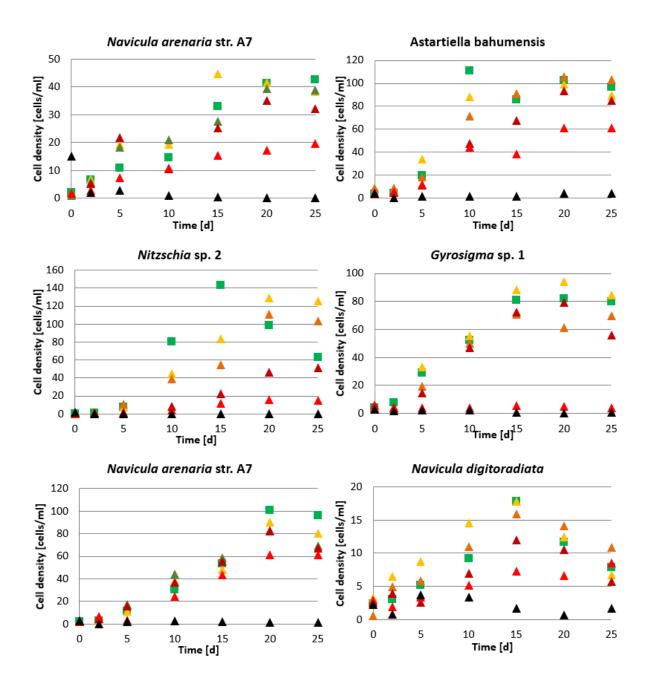




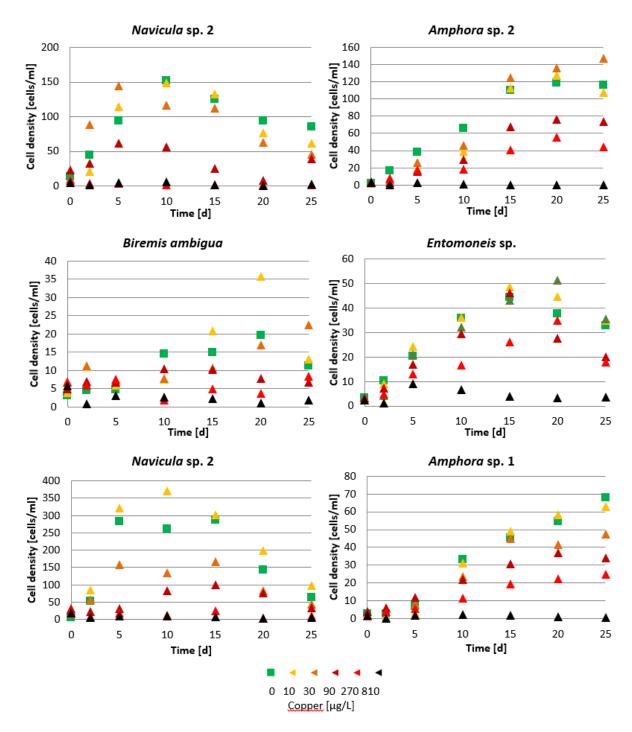


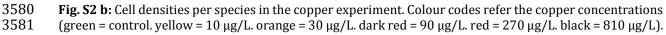
3573Fig. S2 a: Cell densities per species in the atrazine experiment. Colour codes refer the atrazine3574concentrations (green = control. yellow =  $50 \ \mu g/L$ . orange =  $100 \ \mu g/L$ . dark red =  $200 \ \mu g/L$ . red =  $500 \ \mu g/L$ .3575black =  $1000 \ \mu g/L$ ).











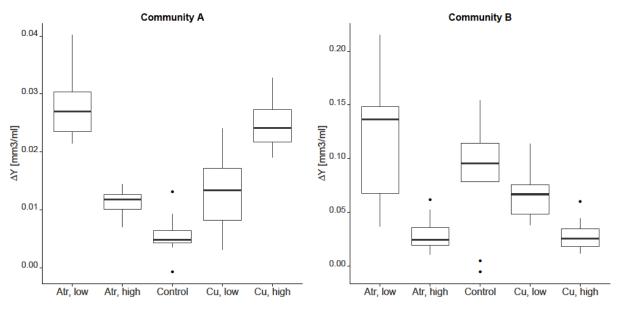
# 3584 Addendum III – Supporting Material Chapter 4

		Acc.	Biovolume	Biovolume	EPS	EPS /
Community	Species	Nr.	total	/ cell	total	biomass
		INI.	[10 <sup>3</sup> mm <sup>3</sup> ]	[µm³]	[mg/ml]	[mg/mm <sup>3</sup> ]
	Nitzschia	DCG	31.46	380	0.25	7.90
	sp. 1	0483	(+- 3.09)	(+- 48)	(+- 0.01)	(+- 0.49)
	Nitzschia	DCG	25.39	517	0.31	12.43
	acicularis	0459	(+- 3.08)	(+- 68)	(+- 0.01)	(+- 1.76)
	Amphora	DCG	18.47	867	0.30	16.65
А	lineolata	0478	(+- 2.82)	(+-247)	(+- 0.01)	(+-3.02)
A	Cylindrotheca		35.68	345	0.32	9.16
	closterium		(+- 2.81)	(+- 60)	(+- 0.03)	(+- 1.68)
	Gyrosigma	DCG	43.52	2081	0.27	6.47
	sp. 2	0491	(13.67)	(+-241)	(+- 0.11)	(+-2.92)
	Navicula	DCG	78.88	5644	0.18	2.37
	arenaria	0487	(18.50)	(+- 723)	(+- 0.01)	(+- 0.49)
	Currentama an 1	DCG	20.62	2147	1.56	75.88
	<i>Gyrosigma</i> sp. 1	0468	(+- 5.49)	(+- 509)	(+- 0.42)	(+- 5.15)
	Entomonoio en	DCG	87.72	10932	0.16	1.93
	<i>Entomoneis</i> sp.	0466	(+- 9.33)	(+- 2743)	(+- 0.02)	(+- 0.30)
	Navicula	DCG	78.88	5644	0.18	2.37
В	arenaria	0487	(18.50)	(+- 723)	(+- 0.01)	(+- 0.49)
D	Amphora	DCG	18.47	867	0.30	16.65
	lineolata	0478	(+-2.82)	(+-247)	(+- 0.01)	(+- 3.02)
	Currentiama en 2	DCG	43.52	2081	0.27	6.47
	<i>Gyrosigma</i> sp. 2	0491	(13.67)	(+-241)	(+- 0.11)	(+- 2.92)
	Navicula	DCG	36.32	5738	1.36	39.19
	digitoradiata	0490	(+- 6.86)	(+- 528)	(+- 0.31)	(+- 14.76)

**Table S1:** Listing of the two diatom communities used in the experiments. From left to right are indicted:
 species name, strain accession number ('Acc. Nr. ') in the diatom culture collection (BCCM/DCG) of the
 Belgian Coordinated Collection of Micro-organisms (http://bccm.belspo.be), biovolume (total and per cell)
 and EPS production (total and per biomass) in monoculture (all n=3).

Substance	Theoretical concentration [µg/l]	Measured Concentration [µg/l]
Control	<1	<2.6
Atrazine	200	160
Atrazine	500	542
Copper	200	271
Copper	500	508

**Table S2:** Experimental atrazine and copper treatments with indication of theoretical and verified atrazine and copper concentrations as determined by GC-MS and graphite furnace AAS respectively (Limits of quantification : < 1,0 μg/L atrazine and 2.6 μg/L copper for 10 ml samples).



**Fig. S1:** The net biodiversity effect in two benthic diatom communities (A and B) exposed to low and high 3602 (200 and 500  $\mu$ g/L) concentrations of copper and atrazine.  $\Delta$ Y is the difference between the biomass yield 3603 observed in diatom communities and the yield expected from the individual species' monocultures.

Com	Stress	Slope	s.e.	Т	Р	Mod	AIC	Log lik.	Validity	LR	P LR
	Atr, low	0.0038	0.0005	6.96	<0.000	1					
	Atr, high	0.0016	0.0005	3.06	0.0040	1	-405	208	Yes		
	Cu, low	0.0012	0.0005	2.15	0.0373		-405	200	Tes		
А	Cu, high	0.0043	0.0005	7.98	<0.0002	1				7 2 2	0 1 2 4 4
A	Atr, low	0.0038	0.0005	8.04	<0.000	1				7.23	0.1244
	Atr, high	0.0017	0.0055	3.42	0.0014	2	402	212	Yes		
	Cu, low	0.0012	0.0006	1.88	0.0667	Z	-403	212	res		
	Cu, high	0.0043	0.0003	12.00	<0.0002	1					
	Atr, low	-0.0072	0.0016	-4.55	<0.000	1					
	Atr, high	-0.0125	0.0016	-7.88	<0.0002	<b>1</b> 1	1 010	165	Yes		
	Cu, low	-0.0215	0.0016	-13.50	<0.000	1	-319	165	Tes		
В	Cu, high	-0.0134	0.0016	-8.41	<0.000	1				15.28	0.0042
D	Atr, low	-0.0072	0.0219	-3.31	0.0019	)				13.20	0.0042
	Atr, high	-0.0125	0.0147	-8.55	<0.0002	1 2	226	172	Voo		
	Cu, low	-0.0215	0.0167	-12.84	<0.0002	1	-326	173	Yes		
	Cu, high	-0.0134	0.0148	-9.01	<0.000	1					

Table S3: Results of generalized least squared models predicting changes in the dominance effect under
 copper and atrazine stress compared to control conditions. 'Com' denotes for which diatom community (A
 or B) the models were fitted. 'Stress' denotes which type of stress diatoms were exposed to: high or low
 (200 or 500 µg/L) copper and atrazine. 'Slope' indicates the relation between the dominance effect and

treatment type, i.e. the difference of the dominance effect (in mm<sup>3</sup>/ml) at each stress level compared to the control. 's.e. ' is the standard error on the estimated slopes. 'T' and 'P' denote the t- and p-values, bold values are statistically significant. 'Mod' indicates if the model was fitted without (Model 1) or with (Model 2) a variance structure. 'AIC' is the Akaike information criterion, 'Log lik' the log-likelihood. 'Validity' denotes if residuals were homogeneous and normally distributed ('yes') or not ('no'). If 'no', models were refitted ('Model 2') with a variance structure allowing the residuals to change according to treatment type. 'LR' is the likelihood ratio of model 1 vs. model 2, P LR the corresponding p-value.

3618

Com	Stress	Slope	s.e.	Т	Р	Mod	AIC	Log lik.	Validity	LR	P LR
	Atr, low	0.0191	0.0022	8.67	<0.0001						
	Atr, high	0.0040	0.0022	1.80	0.0799	1	-293	152 Yes	Voc		
	Cu, low	0.0060	0.0022	2.73	0.0094	1	1 275		Tes		
А	Cu, high	0.0148	0.0022	6.73	<0.0001					- 9.31	0.0537
A	Atr, low	0.0191	0.0024	7.83	<0.0001					9.51	0.0337
	Atr, high	0.0040	0.0014	2.92	0.0057	2	204	157	Voc		
	Cu, low	0.0060	0.0021	2.88	0.0064	Z	2 -294	157	Yes		
	Cu, high	0.0148	0.0019	7.64	<0.0001						
	Atr, low	0.0390	0.0178	2.19	0.0300						
	Atr, high	-0.0417	0.0178	-2.34	0.0242	1	-125	25 69	No		
	Cu, low	0.0029	0.0178	0.16	0.8707	1	-125	09	NU		
В	Cu, high	-0.0421	0.0178	-2.37	0.0229					- 21.91	0 0002
Б	Atr, low	0.0390	0.0258	1.51	0.1392		2 -139			21.91	0.0002
	Atr, high	-0.0417	0.0177	-2.36	0.0234	2		79	Voc		
	Cu, low	0.0029	0.0184	0.16	0.8749	2		79	Yes		
	Cu, high	-0.0421	0.0175	-2.40	0.0210						

3619

3620 Table S4: Results of generalized least squared models predicting changes in the complementarity effect 3621 under copper and atrazine stress compared to control conditions. 'Com' denotes for which diatom 3622 community (A or B) the models were fitted. 'Stress' denotes which type of stress diatoms were exposed to: high or low (200 or 500  $\mu$ g/L) copper and atrazine. 'Slope' indicates the relation between the 3623 3624 complementarity effect and treatment type, i.e. the difference of the complementarity effect (in mm<sup>3</sup>/ml) at each stress level compared to the control. 's.e. ' is the standard error on the estimated slopes. 'T' and 'P' 3625 3626 denote the t- and p-values, bold values are statistically significant. 'Mod' indicates if the model was fitted 3627 without (Model 1) or with (Model 2) a variance structure. 'AIC' is the Akaike information criterion, 'Log lik' the log-likelihood. 'Validity' denotes if residuals were homogeneous and normally distributed ('yes') or not 3628 3629 ('no'). If 'no', models were refitted ('Model 2') with a variance structure allowing the residuals to change 3630 according to treatment type. 'LR' is the likelihood ratio of model 1 vs. model 2, P LR the corresponding p-3631 value.

Com	Effect	Stress	Slope	s.e.	Т	Р	Mod	AIC	Log lik.	Validity	LR	P LR
		Atr, low	0.0189	0.0026	7.08	<0.0001						
		Atr, high	0.0090	0.0026	3.27	0.0022	1	-278	145	Ves		
		Cu, low	0.0072	0.0026	2.72	0.0096	1	270	115	105		
	TIC	Cu, high	0.0141	0.0026	5.32	< 0.0001					- 3.99	0.4076
	110	Atr, low	0.0189	0.0027	6.99	<0.0001					0.77	0.1070
		Atr, high	0.0090	0.0021	4.18	0.0002	2	-274	147	Yes		
		Cu, low	0.0072	0.0021	3.44	0.0014	-	271	117	100		
A		Cu, high	0.0141	0.0025	5.68	<0.0001						
		Atr, low	0.0004	0.0011	0.34	0.7318						
		Atr, high	-0.0047	0.0011	-4.21	0.0001	1	-349	180	No		
		Cu, low	-0.0012	0.0011	-1.06	0.2964	1	517	100	No		
	TDC	Cu, high	0.0008	0.0011	0.68	0.5012					32.73	< 0.000
	IDC	Atr, low	0.0004	0.0006	0.69	0.4907				Yes		<0.0001
		Atr, high	-0.0047	0.0013	-3.55	0.0010	2	-372 19	196			
		Cu, low	-0.0012	0.0004	-2.78	0.0082	2	572	170			
		Cu, high	0.0008	0.0010	0.78	0.4402						
		Atr, low	0.0391	0.0292	1.34	0.1885						0.0007
		Atr, high	-0.018	0.0292	-0.60	0.5529	1	-86	49	Yes	10.16	
		Cu, low	0.1567	0.0292	5.36	<0.0001	I	-00	49			
	TIC	Cu, high	0.0311	0.0292	1.06	0.2938						
	ПС	Atr, low	0.0391	0.0184	2.12	0.0400					19.10	0.0007
		Atr, high	-0.018	0.0151	-1.16	0.2539	2	07	50	Vee		
		Cu, low	0.1567	0.0337	4.64	<0.0001	Z	-97	59	Yes		
D		Cu, high	0.0311	0.0306	1.02	0.3150						
В		Atr, low	-0.0002	0.0233	-0.01	0.9940					-	
		Atr, high	-0.0242	0.0233	-1.04	0.3063			-			
		Cu, low	-0.1538	0.0233	-6.60	<0.0001	1	-104	.04 58	No		
		Cu, high	-0.0732	0.0233	-3.14	0.0032						
	TDC	Atr, low	-0.0002	0.0087	-0.02	0.9837					51.18	<0.000
	I (	Atr, high	-0.0242	0.0047	-5.18	<0.0001	2	-147	84	Yes		.8 <0.000
		Cu, low	-0.1538	0.0279	-5.52	< 0.0001		<b>I</b> I /	4/ 04	+ Yes		
		Cu, high	-0.1338	0.0279								
		, 0	-0.0/32	0.0230	-3.18	0.0028						

**Table S5:** Results of generalized least squared models predicting changes in two-way complementarity3637(TIC) and one-way complementarity (TDC) under copper and atrazine stress compared to control3638conditions. 'Com' denotes for which diatom community (A or B) the models were fitted. 'Effect' indicates3639the complementarity effect (TIC or TDC). 'Stress' denotes which type of stress diatoms were exposed to:3640high or low (200 or 500 µg/L) copper and atrazine. 'Slope' indicates the relation between the respective3641complementarity effect and treatment type, i.e. the difference of the complementarity effect (in mm³/ml) at

ach stress level compared to the control. 's.e. ' is the standard error on the estimated slopes. 'T' and 'P'
denote the t- and p-values, bold values are statistically significant. 'Mod' indicates if the model was fitted
without (Model 1) or with (Model 2) a variance structure. 'AIC' is the Akaike information criterion, 'Log lik'
the log-likelihood. 'Validity' denotes if residuals were homogeneous and normally distributed ('yes') or not
('no'). If 'no', models were refitted (('Model 2') with a variance structure allowing the residuals to change
according to treatment type. 'LR' is the likelihood ratio of model 1 vs. model 2, P LR the corresponding pvalue.

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Com	Stress	Slope	s.e.	Т	Р	Mod	AIC AIC	Log lik.	Validity	LR	P LR
	Atr	1.2e-3	3.7e-4	3.16	0.0158	1	-44	25	Yes	0.43	0.81
۸	A	9.1e-4	3.3e-4	2.76	0.0282	2	-40	25	Yes	0.43	0.01
A	Cu	-2.6e-4	4.2e-4	-0.64	0.5452	1	-36	21	Yes	3.18	0.20
	Cu	1.1e-4	1.9e-4	0.55	0.5964	2	-37	23	Yes	5.10	0.20
В	Atr	1.4e-3	5.1e-3	0.28	0.7884	1	-14	10	Yes	5.04	0.08
Б	Au	-6.5e-3	2.2e-3	3.03	0.0192	2	-15	12	Yes	5.04	0.00
	Cu	1.2e-2	3.6e-3	3.96	0.0115	1	-5.0	5.5	Yes	3.66	0.16
	Cu	2.1e-2	2.7e-3	7.59	0.0001	2	-6.0	8.0	Yes	3.00	0.10

3652

3653 Table S6: Results of generalized least squared models predicting changes in two-way complementarity 3654 (TIC) from diatom EPS production. 'Com' denotes for which diatom community (A or B) the models were 3655 fitted. 'Stress' denotes which type of stress diatoms were exposed to: copper (Cu) and atrazine (Atr). Models 3656 were fitted over the three concentrations of each stressor (control, 200 and 500 µg/L). 'Slope' indicates the 3657 relation between EPS production and TIC. 's.e. ' is the standard error on the estimated slopes. 'T' and 'P' 3658 denote the t- and p-values, bold values are statistically significant. 'Mod' indicates if the model was fitted 3659 without (Model 1) or with (Model 2) a variance structure. 'AIC' is the Akaike information criterion, 'Log lik' 3660 the log-likelihood. 'Validity' denotes if residuals were homogeneous and normally distributed ('yes') or not ('no'). If 'no', models were refitted ('Model 2') with a variance structure allowing the residuals to change 3661 3662 according to treatment type. 'LR' is the likelihood ratio of model 1 vs. model 2, P LR the corresponding p-3663 value.

Com	Stress	Pred	Slope	s.e.	Т	P Mo	d	AIC	Log lik.	Validity	LR	P LR
		Tal	-0.27	0.21	-1.30	0.2240	1	20.0	-7.0	Yes	-0.01	0.05
	A +	Tol	-0.27	0.21	-1.30	0.2213	2	22.0	-7.0	Yes	< 0.01	0.95
	Atr	Viald	-0.23	0.16	-1.38	0.1983	1	20.3	-7.2	No	0.01	0.02
		Yield	-0.23	0.16	-1.39	0.1937	2	-22.3	-7.1	No	0.01	0.92
А		т.l	-0.19	0.25	-0.78	0.4557	1	29.2	-11.6	Yes	0.40	0 5 2
	Con	Tol	-0.21	0.26	-0.79	0.4501	2	30.9	-11.4	Yes	0.40	0.53
	Cu	X7: 1.1	-0.25	0.23	-1.10	0.2982	1	28.8	-11.4	Yes	0.4.4	0.54
	Yield	Yield	-0.27	0.24 -1.13 0.2841 2 30.4 -1	-11.2	Yes	0.44	0.51				
		m 1	-0.41	0.31	-1.33	0.2107	1	21.2	-7.6	Yes	0.4.0	0.66
	<b>A</b> .	Tol	-0.44	0.30	-1.47	0.1735	2	23.0	-7.5	Yes	0.19	0.66
	Atr		-0.21	0.23	-0.89	0.3960	1	22.8	-8.4	Yes	0.4.4	0.75
P		Yield	-0.22	0.23	-0.97	0.3540	2	24.7	-8.3	Yes	0.11	0.75
В		m 1	-0.78	0.21	-3.80	0.0035	1	19.3	-6.9	Yes	0.04	0.65
	Car	Tol	-0.80	0.20	-3.99	0.0026	2	21.2	-6.6	Yes	0.21	0.65
	Cu Yield	-0.54	0.18	-2.99	0.0136	1	22.4	-8.2	Yes			
		Yield	-0.55	0.18	-3.07	0.0118	2	24.3	-8.2	Yes	0.08	0.77

3667 Table S7: Results of generalized least squared models predicting the deviation of observed relative yield 3668 from observed frequency of diatoms grown in atrazine- and copper-exposed communities from the species' 3669 monoculture yields and stress tolerance. 'Com' denotes for which diatom community (A or B) the models 3670 were fitted. 'Stress' denotes which type of stress diatoms were exposed to: copper (Cu) and atrazine (Atr). Models were fitted over the two concentrations of each stressor (200 and 500 µg/L). 'Pred' denotes the 3671 3672 predictor: species monoculture yield ('Yield') or stress tolerance ('Tol'). 'Slope' indicates the relation 3673 between the deviation of the species' observed yield from their observed frequency and their monoculture 3674 yield and stress tolerance. 's.e.' is the standard error on the estimated slopes. 'T' and 'P' denote the t- and 3675 p-values, bold values are statistically significant. 'Mod' indicates if the model was fitted without (Model 1) 3676 or with (Model 2) a variance structure. 'AIC' is the Akaike information criterion, 'Log lik' the log-likelihood. 3677 'Validity' denotes if residuals were homogeneous and normally distributed ('yes') or not ('no'). If 'no', 3678 models were refitted ('Model 2') with a variance structure allowing the residuals to change according to treatment type. 'LR' is the likelihood ratio of model 1 vs. model 2, P LR the corresponding p-value. 3679

## 3681 Addendum IV – Supporting Material Chapter 5

3682

	1.00	Biovolume	Biovolume	EPS	EPS /	EFA	EFA /
Species	Acc. Nr.	total	/ cell	total	biomass	total	biomass
_	INT.	[10 <sup>3</sup> mm <sup>3</sup> ]	[µm³]	[mg/ml]	[mg/mm <sup>3</sup> ]	[µg/ml]	[µg / mm <sup>3</sup> ]
Nitzschia	DCG	31.46	380.34	0.25	7.90	5.80	184.47
sp.	0483	(+- 3.09)	(+- 48)	(+- 0.01)	(+- 0.49)	(+- 0.58)	(+- 4.31)
Nitzschia	DCG	25.39	517.58	0.31	12.43	2.53	98.67
acicularis	0459	(+- 3.08)	(+- 68)	(+- 0.01)	(+- 1.76)	(+- 0.63)	(+- 13.62)
Amphora	DCG	18.47	867.75	0.30	16.65	0.33	18.34
lineolata	0478	(+- 2.82)	(+- 247)	(+- 0.01)	(+- 3.02)	(+- 0.08)	(+- 7.59)
Cylindrotheca		35.68	345.20	0.32	9.16	0.46	13.17
closterium		(+- 2.81)	(+- 60)	(+- 0.03)	(+- 1.68)	(+- 0.17)	(+- 5.24)
Gyrosigma	DCG	43.52	2081.56	0.27	6.47	0.70	16.23
sp.	0491	(+-13.67)	(+- 241)	(+- 0.11)	(+- 2.92)	(+- 0.21)	(+- 3.44)
Navicula	DCG	78.88	5644.09	0.18	2.37	3.29	42.13
arenaria	0487	(+-18.50)	(+- 723)	(+- 0.01)	(+- 0.49)	(+- 0.81)	(+- 8.27)

3683

Table S1: Listing of the six diatom strains used in the experiments. From left to right are indicated: species
 name, strain accession number ('Acc. Nr. ') in the diatom culture collection (BCCM/DCG) of the Belgian
 Coordinated Collection of Micro-organisms (http://bccm.belspo.be), biovolume (total and per cell), EPS and
 EFA production (total and per biovolume) in monoculture (all n=3).

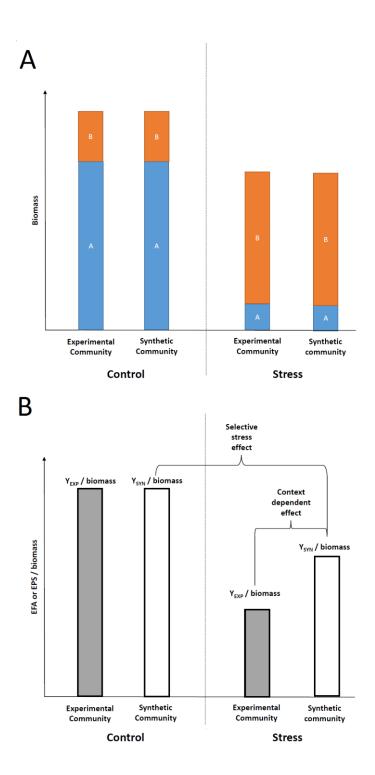
3689

Substance	Theoretical concentration [µg/l]	Measured Concentration [µg/l]
Control	<1	<2.6
Atrazine	200	160
Atrazine	500	542
Copper	200	271
Copper	500	508

3690

**Table S2:** Experimental atrazine and copper treatments with indication of theoretical and verified atrazine and copper concentrations as determined by GC-MS and graphite furnace AAS respectively (Limits of quantification : <  $1,0 \mu$ g/L atrazine and  $2.6 \mu$ g/L copper for 10 ml samples).

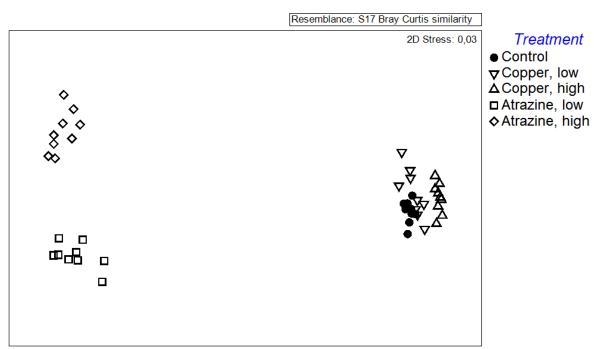
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3697 Fig. S1: Quantification of selective and context-dependent stress effects by comparison of the EFA and EPS 3698 production in experimental and synthetic communities. For the sake of clarity, this example uses a 3699 community with only two species (A and B) and only one stress level. Panel A shows the community 3700 structure of experimental and synthetic communities. In the experimental community, stress induces a 3701 change in biomass and community structure (dominance of species B under stress). The synthetic 3702 community reflects the biomass and community structure observed in the corresponding experimental 3703 community. Panel B shows the EFA or EPS production per unit biomass in the experimental community 3704 (Y<sub>EXP</sub>) and in the synthetic community (Y<sub>SYN</sub>). Y<sub>SYN</sub> is calculated by multiplying each species' EFA or EPS 3705 production in unstressed monocultures with its biomass share in the corresponding experimental 3706 community. Y<sub>SYN</sub> thus reflects the EFA or EPS production expected in a community without stress exposure 3707 and species interactions, but which has the same biomass and community structure as induced by stress. 3708 Selective stress effects are quantified by comparing synthetic communities reflecting the community

3709 structure under stress and under control conditions. Confounding effects of direct stress or changes in 3710 species interactions are eliminated, since Y<sub>SYN</sub> of both communities is calculated from unstressed 3711 monocultures. Context-dependent effects are quantified by comparing experimental and synthetic 3712 communities of the same community structure. Confounding effects of community structure are eliminated, 3713 and differences in Y<sub>EXP</sub> and Y<sub>SYN</sub> are attributed to species interactions or direct stress effects on Y<sub>EXP</sub>. In this example, Y<sub>EXP</sub> is lower under stress than in the control. This is due to both context-dependent stress effects 3714 3715 (Y<sub>EXP</sub> stress < Y<sub>SYN</sub> stress) and selective stress effects (Y<sub>SYN</sub> stress < Y<sub>SYN</sub> control). Here, stress would thus reduce diatom EFA or EPS production by reducing the species' EFA or EPS output, as well as by causing 3716 3717 dominance by an unproductive species.

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Fig. S2: Non-metric multidimensional scaling (Bray-Curtis similarity) of untransformed relative (%) biomass data of communities from unstressed conditions and exposed to low (200  $\mu$ g/L) and high (500 3723  $\mu$ g/L) atrazine and copper concentrations.

3724

Treatments	Dissimilarity	Species	Contribution
	Navicula arenaria		33.25
Copper, low	14.49	Nitzschia acicularis	33.00
		<i>Gyrosigma</i> sp.	10.92
		Navicula arenaria	28.56
Copper, high	18.30	<i>Nitzschia</i> sp.	19.60
		Nitzschia acicularis	16.51
		Cylindrotheca closterium	49.63
Atrazine, low	69.79	Nitzschia acicularis	28.45
		Navicula arenaria	12.71
		Nitzschia acicularis	39.20
Atrazine, high	75.97	Cylindrotheca closterium	25.23
	Γ	Navicula arenaria	24.79

3726 Table S3. Results of dissimilarity percentage (SIMPER) analysis of biomass composition of diatom 3727 communities grown exposed to copper and atrazine compared to control communities. Analysis was run on 3728 the untransformed biomass data. 'Treatments' indicates the treatment whose biomass composition is 3729 compared to the control: low (200  $\mu$ g/L) and high (500  $\mu$ g/L) atrazine and copper. 'Dissimilarity' indicates 3730 the average dissimilarity (in %) of biomass composition of the respective treatment and the control. 'Species' indicates the three species contributing most to dissimilarities with the control biomass 3731 3732 composition. 'Contribution' indicates the average contribution (in %) of each species to dissimilarity from 3733 the control biomass composition.

3734

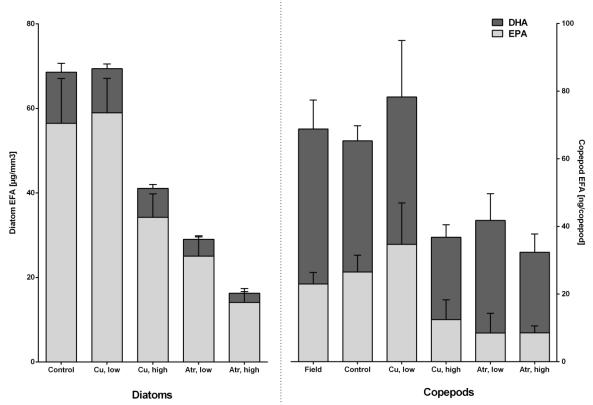
### 3735

Treatment	Similarity	Species	Mean	Contribution
		Nitzschia acicularis	20.01	48.02
Control	91.52	Navicula arenaria	15.17	32.56
		<i>Nitzschia</i> sp.	2.78	5.93
		Nitzschia acicularis	18.30	49.25
Copper, low	84.07	Navicula arenaria	12.69	33.26
		<i>Nitzschia</i> sp.	2.70	7.47
Connor high	90.06	Nitzschia acicularis	20.96	62.52
Copper, high	90.06	Navicula arenaria	11.62	33.10
		Cylindrotheca closterium	36.15	76.27
Atrazine, low	88.32	Navicula arenaria	7.44	9.85
		<i>Gyrosigma</i> sp.	4.78	3.78
		Cylindrotheca closterium	15.68	74.22
Atrazine, high	86.27	<i>Gyrosigma</i> sp.	2.76	11.90
		Navicula arenaria	2.46	9.06

3736

3737 Table S4: Results of similarity percentage (SIMPER) analysis of biomass composition of diatom 3738 communities grown under unstressed conditions as well as copper and atrazine exposure. Analysis was run 3739 on the untransformed biomass data. 'Treatment' indicates the growth conditions: control, low (200 µg/L) 3740 and high (500 µg/L) atrazine and copper. 'Similarity' indicates the average similarity (in %) of biomass 3741 composition within the nine replicates of each treatment. 'Species' indicates the species contributing most 3742 to similarities within treatments (only species with a mean biomass >  $2 \times 10^3 \mu m^3$  are reported). 'Mean' 3743 indicates the mean biomass (in 10<sup>3</sup> µm<sup>3</sup>) of species within treatments. 'Contribution' indicates the 3744 contribution (in %) of each species to similarities within treatments.

3745



3747DiatomsCopepods3748Fig. S3: EPA and DHA content of diatom communities exposed to low (200 μg/L) and high (500 μg/L)3749atrazine ('Atr') and copper ('Cu') concentrations, and EPA and DHA content of *Microarthridion littorale*3750collected in the field and after 10d of feeding on the diatom diet that was subject to the stressors (see left3751part of the figure for corresponding abbreviations of the tractments)

3751 part of the figure for corresponding abbreviations of the treatments).

### 3753 Addendum V – Acute toxicity of copper to harpacticoid copepods

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Addendum V reports the results of single-species toxicity tests, which measured the tolerance offive harpacticoid copepod species to direct copper exposure.

3757 Harpacticoid copepods were collected from sediments at the Paulina intertidal flat (SW 3758 Netherlands, 51° 21'N, 3°43'E). Five harpacticoid copepod species (Microathridion littorale, 3759 Stenhelia sp., Platychelipus littoralis, Nannopus palustris and one species of the family 3760 Ectinosomatidae) were extracted alive from the sediment using a mixed technique of sediment 3761 decantation and extraction via white light attraction. Adult specimens were randomly collected 3762 with a glass pasteur pipette using a Wild M5 binocular. Copepods were washed 3 times over a 38 3763 µm sieve and placed in glass jars with filtered and autoclaved natural seawater overnight in order 3764 to empty their intestines prior to the start of the experiment.

3765 Copepod sensitivity to direct copper exposure was quantified by exposing the five harpacticoid 3766 species collected at the sampling site for 48h to a concentration gradient of 0 to 2430  $\mu$ g/L copper 3767 (Table S1). The experiment was conducted in glass jars containing 100 ml of filtered and 3768 autoclaved seawater in a climate room at 15±1 °C with a 12:12 h light:dark cycle and 40 to 50 3769  $\mu$ mol photons m<sup>-2</sup> s<sup>-1</sup>. During the experiment, harpacticoids were offered 0.5 mm<sup>3</sup> of an equal mix 3770 of the six diatom species used as food source in Chapter 5 (*Nitzschia* sp., *N. acicularis, C. closterium*, 3771 *A. lineolata, Gyrosigma* sp., *N. arenaria*).

Cu	Replicates (copepod individuals per replicate)								
[µg/l]	Microarthridion littorale	<i>Stenhelia</i> sp.	Nannopus palustris	Ectinosomatidae sp.	Platychelipus littoralis				
0	5 (15)	5 (15)	5 (15)	5 (15)	5 (30)				
10	5 (15)				5 (30)				
30	5 (15)	5 (15)	5 (15)	5 (15)	5 (30)				
90	5 (15)				5 (30)				
270	5 (15)	5 (15)	5 (15)	5 (15)	5 (30)				
810	5 (15)	5 (15)	5 (15)		5 (30)				
2430	5 (15)	5 (15)	5 (15)	5 (15)	5 (30)				

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Table S1: Experimental copper treatments with indication of the number of replicates per treatment and
 the number of individuals per replicate (in brackets) for each copepod species (*Microarthridion littorale, Stenhelia* sp., *Nannopus palustris*, Ectinosomatidae sp., *Platychelipus littoralis*)

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3777 Copepod sensitivity to direct copper exposure was quantified as the 48-h median lethal 3778 concentration (LC50), i.e. the effective copper concentration inducing a 50% mortality of each harpacticoid species. LC50 values were calculated by means of a three-parameter log-logistic
model (Equation 1) and a three-parameter Weibull model in two different parameterisations
(Equations 2 and 3) as described in Ritz & Streibig (2005).

3782	$Y = d / (1 + \exp\{b[\log x - \log e]\})$	[1]
3782	$Y = d / (1 + \exp\{b[\log x - \log e]\})$	[1

3783 
$$Y = d(1 - \exp[b(\log x - \log e)])$$
 [2]

3784  $Y = d(\exp\{-\exp[b(\log x - e)]\})$ 

3785 *Y* represents the response variable (surviving harpacticoid individuals), *b* the relative slope of the
3786 curve, *d* the upper limit, *e* the inflection point, and *x* the copper concentration. Dose-response
3787 models and LC50 values were computed in R 3.0.1. using RStudio (R Development Core Team
3788 2016) and the package drc (Ritz and Streibig 2005).

[3]

3789

The five harpacticoid copepod species differed in their sensitivity to direct copper exposure, with LC50 values ranging from 387  $\mu$ g/L (*N. palustris*) to 1475  $\mu$ g/L (*Stenhelia* sp.). *M. littorale*, being the most abundant copepod species at the sampling site representing ~ 90% of all collected individuals, was the second most copper-sensitive species with an LC50 of 546  $\mu$ g/L (Table S2).

Species	LC50	Surviving individuals per treatment (µg/L Cu)						
Species		Control	10	30	90	270	810	2430
Microarthridion littorale	546.43 (+-78.38)	65	67	68	66	48	18	14
Stenhelia sp	1474.60 (+-235.58)	71		70		57	55	14
Nannopus palustris	386.83 (+-160.89)	67		67		40	23	0
Ectinosomatidae sp.	696.88 (+-188.42)	66		50		48	33	2
Platychelipus littoralis	879.28 (+-260.09)	134	133	107	116	115	67	12

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Table S2: Copper 48h-LC50 values obtained for five harpacticoid copepod species (standard deviations are
 indicated in brackets). Each treatment had five replicates, with 15 individuals (30 for *Platychelipus littoralis*)
 harpacticoid individuals per replicate. Surviving individuals are indicated as sum of the five replicates.

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