

- 1 ExMarine: An experimental field mesocosm system to study
- 2 multiple-stressor impacts on rock pool biodiversity

- 4 Author names and affiliations
- 5 Jan N. Macher<sup>1\*</sup>,
- <sup>1</sup> Aquatic Ecosystem Research, University Duisburg-Essen, Universitätsstraße 5, 45141
- 7 Essen, Germany
- 8 \*Corresponding author: jan.macher@uni-due.de
- 9 Keywords: Mesocosm, Tide pool, Experimental setup, Experimental design,
- 10 Ecological experiment, Multiple stressors, Shore ecosystem

11

12

13

14

15

16

17

18

19

20

21

22

23

24

25

26

#### Abstract

Biodiversity loss due to increasing anthropogenic activities is one of the biggest threats to humanity. Understanding the impacts of multiple-stressors on ecosystems and biodiversity is therefore an urgent task. Shore ecosystems are especially valuable, as they harbour a high biodiversity and provide important ecosystems services. Until now, experimental approaches addressing multiple-stressor impacts on these ecosystems have been rare and mostly run with a limited number of replicates and under non-natural conditions. Here, an experimental field mesocosm system that allows studying multiple-stressor impacts on rock pool biodiversity is proposed. The *ExMarine* mesocosm system is composed of 64 experimental rock pool mesocosms in a fully randomised block design, which allows studying multiple-stressor impacts under highly standardised conditions. Water is taken directly from the sea, allowing biota to immigrate and emigrate freely. Water flow into the mesocosms can be regulated and it is possible to simulate disturbance through waves during high tide. The system can help to understand the impacts of multiple stressors on biodiversity, to monitor ecosystem health and to plan measures preventing the further loss of biodiversity.



## 27 Introduction

2829

30

31

32

33

34

35

36

37

38

39

40

41 42

43

44

45

46

47

48

49

50

51

52

53

54

55

56

57

58

59

60

The ongoing loss of biodiversity is one of the biggest threats to humanity (Rockström et al. 2009). As biodiversity provides food resources (Bommarco et al. 2013) and important ecosystem services such as water purification (see e.g. Edwards & Abivardi 1998; Vörösmarty et al. 2010) and carbon pulldown (see e.g. Hügler & Sievert 2011; Gilroy et al. 2014), societies worldwide rely on intact ecosystems and their biodiversity. However, anthropogenic activities such as intensified land use, industrialisation and overexploitation of natural resources have been shown to impact all levels of biodiversity from genes to ecosystems (Steffen et al. 2015). These multiple anthropogenic stressors can interact in different ways, sometimes leading to "ecological surprises" and impacting ecosystems and biodiversity in unexpected ways (Piggott et al. 2015a; Jackson et al. 2016). Therefore, an indepth understanding of complex multiple-stressor interactions is urgently needed in order to prevent the further loss of biodiversity and ecosystems functions. To gain this understanding, studying ecosystems that harbour a high biodiversity is especially promising. Studies have addressed the impacts of multiple stressors on different ecosystems, from terrestrial (e.g. Luers et al. 2003; Davalos et al. 2014) to freshwater (e.g. Piggott et al. 2015b; Macher et al. 2016) and marine (e.g. Ban et al. 2014; Ellis et al. 2015). One marine ecosystem that is especially suitable for studying multiple-stressor effects is the rocky shore. Rocky shores are highly dynamic areas, affected by both sea and land, and harbour a high number of habitats and species communities along a relatively narrow strip (Levinton 1995). In addition, rocky shores are of significant economic value in many parts of the world, as a large number of edible species (such as crabs and seaweeds) are harvested in this ecosystem (e.g. Duran & Castilla 1989). One feature of rocky shores is the formation of rock or tide pools in the intertidal zone. The intertidal zone is mostly submerged at high tides and during storms, but can be exposed to wind, freshwater influx and extreme temperatures during low tide. Due to these conditions, rock pools are considered extreme habitats (Knox 2000). Rock pools have been studied intensively due to their accessibility, their diverse and ecologically interesting species communities (e.g. Connell 1961, Oksanen et al. 2002; Vanschoenwinkel et al. 2007; Brendonck et al. 2015) and due to the fact that they can be used as natural mesocosms in experimental studies (e.g. Dethier 1984, Romanuk et al. 2009). Therefore, they seem perfectly suitable for studies on multiple-stressor impacts on biodiversity and ecosystem functions. However, experimental studies on rock pools to date have been mostly conducted either in natural rock pools, which vary in size, depth, position relative to the shore and which therefore do not allow conducting experiments under strictly controlled conditions (e.g.

# Peer | Preprints

Dethier 1984; Romanuk et al. 2009) or in mostly artificial pools, which suffer from drawbacks such as closed community structure or lack of water influx from the sea (e.g. Sloman et al. 2008). Some studies have used intermediate designs, e.g. by building artificial pools close to the shoreline that are allowed to flood regularly (e.g. Nielsen 2001, Romanuk 2009), but that do not allow a manipulation of physicochemical parameters or addition of stressors throughout the time of the experiment. In all studies, the number of replicates has been rather low. None of the existing systems allows manipulating physicochemical parameters in a highly replicated mesocosms system under natural conditions. Here, an experimental system that allows studying the impacts of multiple stressors on rock pools under highly realistic and standardised conditions and with a high number of replicates in a fully randomised block design is proposed. The system is based on the *ExStream* field mesocosm system, which was first used by Lange et al. (2011) to study freshwater species communities under multiple stressors and which has since been successfully used in several large experiments (e.g. Piggott et al. 2012, Elbrecht et al. 2016). The here-proposed system is called *ExMarine* (Experimental Marine Rock Pool Mesocosm System).

#### **Material and Methods**

As described in Lange et al. (2011) and Piggott et al. (2015b), the experimental system consists of mesocosms that are placed on a two-storey scaffold next to the studied habitat, therefore allowing studying the ecosystem of interest under natural conditions. For the ExMarine system, the scaffold is set up next to an area of shore with rock pools. A pump inlet with filters to prevent the intake of coarse material is placed in the intertidal zone. The pump transports water to the scaffold during high tides only in order to simulate the fluctuating supply of water during high and low tides. Pumps can either be switched on and off manually or activated via a programmed timer. If required, the pump can also run and supply the mesocosms with water at all times if placed in deeper water below the intertidal zone. It is also possible to run the pumps at longer intervals to allow desiccation of the rock pool mesocosms. The water pumped from the seas is distributed into eight header tanks placed on the second-storey platform of the scaffold. Each header tank then distributes the water into eight rock pool mesocosms via gravity (Figure 1). The flow rate can be controlled via valves, which are installed upstream of every channel. When water is supplied to the mesocosms during high tide, the excess water leaves the mesocosms via an outlet. During low tides, when the pump is not working, the water level in the mesocosm falls below the outflow level. In

# Peer | Preprints

95

96

97

98

99

100

101

102

103

104

105

106

107

108

109

110

111

112

113

114

115

total, the ExMarine system consists of 64 rock pool mesocosms. Prior to an experiment, the mesocosms can be filled with substratum and can be stocked with biota from the surrounding rock pools. Throughout the experiment, small biota (mainly microbes, algae and invertebrates) get pumped into the experimental system directly from the sea together with the water for the mesocosms. They can then settle in the mesocosms or leave them through the outlet during simulated high tide phases. The ExMarine system can be built in two different ways. One possibility is to build it so that water is directly pumped into the header tanks, from where it is distributed to the mesocosms via gravity (Figure 1, a + b). This setup does allow simulation of high tide and low tide phases, but not of varying water influx during high tide, which is typical for rock pools on the intertidal zone. Therefore, it is also possible to build the setup with water scoops, which are located next to the header tanks (Figure 1, c + d). During high tide, water is pumped into the water scoops, gradually filling them. Thereby, the water scoops become heavier and finally tilt, releasing the water into the header tanks in a squall. From the header tanks, the squall is distributed to the mesocosms, simulating the varying influx of water and the turbulences caused by waves. After the completion of the experimental phase, the community in each channel is assessed either by morphological identification or molecular methods such as barcoding (Hebert et al. 2003), metabarcoding (see e.g. Taberlet et al. 2012; Leray & Knowlton 2015), metagenomics (see e.g. Riesenfeld et al. 2004; Crampton-Platt et al. 2015) or metatransciptomics (see e.g. Shi et al. 2009; Marchetti et al. 2012) in order to assess the impacts of the stressors on community structure, species diversity, genetic diversity and ecosystem functions.

**Figure 1:** *ExMarine* rock pool mesocosm system. a) Top view; system with water scoops installed. b) Side view; system with water scoops installed. c) Top view; system without water scoops. d) Side view; system without water scoops.

117

118



## 119 Discussion

120

121

122

123

124

125

126

127

128

129

130

131

132

133

134

135

136

137

138

139

140

141

142

143

144

145

146

147

148

149

150

151

The here-proposed ExMarine field mesocosm system allows studying the impacts of multiple stressors on biodiversity in rock pools, bringing together the statistical power of highly replicated experimental approaches and the benefits of a real-world setting with free immigration and emigration of biota. This approach has been shown to be highly effective in freshwater ecosystems (e.g. Piggott et al. 2015b) and is novel for the study of biodiversity in shore ecosystems. Stressors can be applied to channels via drippers connected to pumps (e.g. nutrients, see e.g. Elbrecht et al. 2016) or added manually (e.g. sediment). Existing experimental rock pool mesocosm systems mostly suffer from drawbacks such as low number of replicates, non-realistic settings, closed communities or (in case of artificial pools build on the shore) the problem that stressors cannot be applied throughout the whole experimental phase due to waves washing them out. In contrast, the ExMarine mesocosm systems offers a solution to these drawbacks. Due to the highly replicated and randomised block design (eight blocks with eight channels each), three stressors can be tested alone and in combination. Together with a control treatment, each stressor and stressor combination can be tested in eight replicates (one per block). Two stressors plus control can be tested in 16 replicates each (two per block). The system can also be enhanced and run with 128 or more channels, or downscaled and run with 32 channels. The great advantage of this experimental system over existing systems is the fact that it is run with water directly from the natural habitat, that specimens can colonise the mesocosms, and that they can leave the experimental system should the conditions in the mesocosms be unfavourable. It has to be acknowledged, however, that pumping of water directly from the sea prevents larger organisms from entering the mesocosms, making it impossible for large predators such as fish to impact on the community in the mesocosms. Also, disturbance through waves might be stronger in real rock pools. However, the experiments using the ExStream stream mesocosm system have shown that results are meaningful and the community is similar to that in the natural environment. Therefore, it can be concluded that the *ExMarine* rock pool mesocosms work equally well and can give insights into the studied ecosystem, allowing to study the ecosystem under highly realistic conditions. Further, since rock pools are extreme, often shallow and small habitats, the mesocosms are expected to accurately depict this habitat. Therefore, the community found in the mesocosms is expected to closely resemble the one in a real rock pool, a fact that can and should be verified by sampling nearby natural rock pools. Overall, the ExMarine system can help to deepen the understanding of how multiple stressors impact on an important and

- highly diverse ecosystem. The experimental system can be used to monitor ecosystem health
- and the results could be helpful when planning conservation measures.

- 155 References
- Ban, S. S., Graham, N. A., & Connolly, S. R. (2014). Evidence for multiple stressor i nteractions and effects on coral reefs. *Global change biology*, 20(3), 681-697.

158

Bommarco, R., Kleijn, D., & Potts, S. G. (2013). Ecological intensification: harnessing ecosystem services for food security. *Trends in ecology & evolution*, 28(4), 230-238.

161

Brendonck, L., Jocqué, M., Tuytens, K., Timms, B. V., & Vanschoenwinkel, B. (2015).
 Hydrological stability drives both local and regional diversity patterns in rock pool
 metacommunities. *Oikos*, 124(6), 741-749.

165

166 Connell, J. H. (1961). The influence of interspecific competition and other factors on the distribution of the barnacle Chthamalus stellatus. *Ecology*, 42(4), 710-723.

168

169 Crampton-Platt, A., Timmermans, M. J., Gimmel, M. L., Kutty, S. N., Cockerill, T. D., Khen, 170 C. V., & Vogler, A. P. (2015). Soup to tree: the phylogeny of beetles inferred by 171 mitochondrial metagenomics of a Bornean rainforest sample. *Molecular biology and evolution*, 32(9), 2302-2316.

173

Dávalos, A., Nuzzo, V., & Blossey, B. (2014). Demographic responses of rare forest plants to multiple stressors: the role of deer, invasive species and nutrients. *Journal of Ecology*, 102(5), 1222-1233.

177178

Duran, L. R., & Castilla, J. C. (1989). Variation and persistence of the middle rocky intertidal community of central Chile, with and without human harvesting. *Marine Biology*, 103(4), 555-562.

180 181

179

Dethier, M. N. (1984). Disturbance and recovery in intertidal pools: maintenance of mosaic patterns. *Ecological Monographs*, *54*(1), 99-118.

184

Edwards, P. J., & Abivardi, C. (1998). The value of biodiversity: where ecology and economy blend. *Biological Conservation*, 83(3), 239-246.

187

Elbrecht, V., Beermann, A. J., Goessler, G., Neumann, J., Tollrian, R., Wagner, R., ... & Leese, F. (2016). Multiple-stressor effects on stream invertebrates: a mesocosm experiment manipulating nutrients, fine sediment and flow velocity. *Freshwater Biology*.

192193

194

195

Ellis, J. I., Hewitt, J. E., Clark, D., Taiapa, C., Patterson, M., Sinner, J., ... & Thrush, S. F. (2015). Assessing ecological community health in coastal estuarine systems impacted by multiple stressors. *Journal of Experimental Marine Biology and Ecology*, 473, 176-187.

196 197

Gilroy, J. J., Woodcock, P., Edwards, F. A., Wheeler, C., Baptiste, B. L., Uribe, C. A. M., ...
 & Edwards, D. P. (2014). Cheap carbon and biodiversity co-benefits from forest
 regeneration in a hotspot of endemism. *Nature Climate Change*, 4(6), 503-507.

Hebert, P. D., Ratnasingham, S., & de Waard, J. R. (2003). Barcoding animal life:
 cytochrome c oxidase subunit 1 divergences among closely related species.
 *Proceedings of the Royal Society of London B: Biological Sciences*, 270, S96-S99.

Hügler, M., & Sievert, S. M. (2011). Beyond the Calvin cycle: autotrophic carbon fixation in the ocean. *Marine Science*, 3.

Jackson, M. C., Loewen, C. J., Vinebrooke, R. D., & Chimimba, C. T. (2016). Net effects of multiple stressors in freshwater ecosystems: a meta-analysis. *Global change biology*, 22(1), 180-189.

214 Knox, G. A. (2000). The ecology of seashores. CRC press.

Lange, K., Liess, A., Piggott, J. J., Townsend, C. R., & Matthaei, C. D. (2011). Light,
 nutrients and grazing interact to determine stream diatom community composition and
 functional group structure. Freshwater Biology, 56(2), 264-278.

Leray, M., & Knowlton, N. (2015). DNA barcoding and metabarcoding of standardized samples reveal patterns of marine benthic diversity. *Proceedings of the National Academy of Sciences*, 112(7), 2076-2081.

Levinton, J. S. (1995). *Marine biology: function, biodiversity, ecology*). New York: Oxford
 University Press.

Luers, A. L., Lobell, D. B., Sklar, L. S., Addams, C. L., & Matson, P. A. (2003). A method for quantifying vulnerability, applied to the agricultural system of the Yaqui Valley, Mexico. *Global Environmental Change*, 13(4), 255-267.

Macher, J. N., Salis, R. K., Blakemore, K. S., Tollrian, R., Matthaei, C. D., & Leese, F. (2016). Multiple-stressor effects on stream invertebrates: DNA barcoding reveals contrasting responses of cryptic mayfly species. *Ecological Indicators*, 61, 159-169.

Marchetti, A., Schruth, D. M., Durkin, C. A., Parker, M. S., Kodner, R. B., Berthiaume, C. T., ... & Armbrust, E. V. (2012). Comparative metatranscriptomics identifies molecular bases for the physiological responses of phytoplankton to varying iron availability. *Proceedings of the National Academy of Sciences*, 109(6), E317-E325.

O'Brien, K., Leichenko, R., Kelkar, U., Venema, H., Aandahl, G., Tompkins, H., ... & West, J. (2004). Mapping vulnerability to multiple stressors: climate change and globalization in India. *Global environmental change*, 14(4), 303-313.

Oksanen, I., Lohtander, K., Paulsrud, P., & Rikkinen, J. (2002, January). A molecular approach to cyanobacterial diversity in a rock-pool community involving gelatinous lichens and free-living Nostoc colonies. In *Annales Botanici Fennici* (pp. 93-99). Finnish Zoological and Botanical Publishing Board.

Piggott, J. J., Lange, K., Townsend, C. R., & Matthaei, C. D. (2012). Multiple stressors in agricultural streams: a mesocosm study of interactions among raised water temperature, sediment addition and nutrient enrichment. *PloS one*, 7(11), e49873.

255

259

262263

266267

268269

272

277

281

285

289

- Piggott, J. J., Townsend, C. R., & Matthaei, C. D. (2015a). Reconceptualizing synergism and antagonism among multiple stressors. *Ecology and evolution*, *5*(7), 1538-1547.
- Piggott, J. J., Salis, R. K., Lear, G., Townsend, C. R., & Matthaei, C. D. (2015b). Climate
   warming and agricultural stressors interact to determine stream periphyton community
   composition. *Global change biology*, 21(1), 206-222.
- Riesenfeld, C. S., Schloss, P. D., & Handelsman, J. (2004). Metagenomics: genomic analysis of microbial communities. *Annu. Rev. Genet.*, *38*, 525-552.
- Rockström, J., Steffen, W., Noone, K., Persson, Å., Chapin, F. S., Lambin, E. F., ... &
  Nykvist, B. (2009). A safe operating space for humanity. *Nature*, 461(7263), 472-475.
  - Romanuk, T. N., Vogt, R. J., & Kolasa, J. (2009). Ecological realism and mechanisms by which diversity begets stability. *Oikos*, *118*(6), 819-828.
- Shi, Y., Tyson, G. W., & DeLong, E. F. (2009). Metatranscriptomics reveals unique microbial
   small RNAs in the ocean's water column. *Nature*, 459(7244), 266-269.
- Sloman, K. A., Mandic, M., Todgham, A. E., Fangue, N. A., Subrt, P., & Richards, J. G.
   (2008). The response of the tidepool sculpin, Oligocottus maculosus, to hypoxia in laboratory, mesocosm and field environments. *Comparative Biochemistry and Physiology Part A: Molecular & Integrative Physiology*, 149(3), 284-292.
- Steffen, W., Richardson, K., Rockström, J., Cornell, S. E., Fetzer, I., Bennett, E. M., ... &
   Folke, C. (2015). Planetary boundaries: Guiding human development on a changing planet. *Science*, 347(6223), 1259855.
- Taberlet, P., Coissac, E., Pompanon, F., Brochmann, C., & Willerslev, E. (2012). Towards
   next-generation biodiversity assessment using DNA metabarcoding. *Molecular ecology*, 21(8), 2045-2050.
- Vanschoenwinkel, B., De Vries, C., Seaman, M., & Brendonck, L. (2007). The role of metacommunity processes in shaping invertebrate rock pool communities along a dispersal gradient. *Oikos*, *116*(8), 1255-1266.
- Vörösmarty, C. J., McIntyre, P. B., Gessner, M. O., Dudgeon, D., Prusevich, A., Green, P., ...
  & Davies, P. M. (2010). Global threats to human water security and river biodiversity.
  Nature, 467(7315), 555-561.