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1 **A simple and rapid method for separation and isolation of 1 marine algal species from**  
2 **naturally evolved populations**

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29 *Study on mixed microalgal populations*

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32 **ABSTRACT**

33 To improve the study of mixed microalgal populations, three naturally evolved marine microalgal  
34 cultures were subjected to a light crushing mechanical treatment using a silicon spatula coupled  
35 with zymolyase treatment at four concentrations: 5, 10, 20 and 25 U/ml, for 15, 30, 45 and 60 min  
36 before being observed under a microscope. The enzyme concentration of 20 U/ml after 45 min  
37 reduces the size of macroscopic microalgal aggregates and improves the microscopic observation of  
38 the different microalgal species comprising the population. There was no improvement using the  
39 higher enzyme concentration. This paper proposes a new approach to the study of naturally evolved  
40 microalgal populations which is useful for distinguishing the morphology of the different species  
41 present in the population and allowing for the identification by classical keys, and also to obtain a  
42 pure culture from an inoculum of mixed species by using a micromanipulator for cell counting.

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44 **Key words:** mixed microalgal populations, natural evolution, zymolyase, mechanical treatment,  
45 algal EPS

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## 55 **Introduction**

56 Nowadays, microalgae are an interesting field to study and explore, as they allow for the production  
57 of an attractive form of alternative energy from lipids (Demirbas 2011). Due to high lipid  
58 production, marine microalgae have been considered potential sources of renewable energy (Jiang  
59 et al. 2011). The great majority of papers on this topic are about the optimization of uni-algal  
60 culture as regards growth rapidity and the high lipid accumulation in the cells (Duong et al. 2012;  
61 Lim et al. 2012; Xie et al. 2012). The use of mixed microalgae naturally evolved from water bodies  
62 resulted in more cost-effective processes. Few authors have reported about induced or natural mixed  
63 algal populations (Phatarpekar et al. 2000; Mohan et al. 2011; Wahlen et al. 2011).

64 This study points out some of the difficulties encountered in studying naturally evolved mixed  
65 microalgae populations, such as in the use of a spectrophotometer; in fact, the different species  
66 clump together, forming macroscopic aggregates that in some cases float and remain in suspension  
67 while in others they tend to sediment after mixing. The behaviour observed is probably attributable  
68 to the production of extracellular polysaccharides (EPS; Kristensen et al. 2008), which impairs the  
69 easy recognition of the different type of cells as well as the identification by classical keys. This  
70 makes it difficult to obtain pure cultures by micromanipulation and to achieve growth curves for the  
71 populations present in the mixed culture. Different enzymes such as chitinase, lysozyme and  
72 pectinase, coupled with other types of treatments, have been used to break down the algal cell walls  
73 to allow release of the lipid (Gerken et al. 2013). Some authors verified the effect of the treatments  
74 by microscopic observations (Allard et al. 2002; Versteegh & Blokker 2004), some others by  
75 electron microscopy (Jensen & Sicko 1971). No paper reports on the use of enzymatic treatment to  
76 attack the microalgal EPS to disaggregate the clumped cells naturally produced from water bodies.

77 The aim of this work is to propose an enzymatic based procedure that allows disaggregating the

78 macro-aggregates of mixed microalgal populations isolated from the marine environment in order to  
79 facilitate the study of naturally evolved populations.

80

## 81 **Material and methods**

82 Three naturally evolved marine mixed microalgal blooms – indicated as A, B and C – were used.  
83 The seawater was collected during the autumn season from different sites along the coast of Reggio  
84 Calabria (Calabria, South Italy), passed through a 100 µm filter, enriched with Walne medium (1  
85 ml/l) plus vitamins (0.1 ml/l) (Walne 1966) and kept at room temperature and natural photoperiod  
86 conditions.

87 The samples were monitored daily by microscopic observation (Zeiss standard 20 optical  
88 microscope) from the day of sampling to the time the macroscopic microalgal population  
89 developed. Then, the development of the A, B and C blooms was monitored microscopically each  
90 week.

91 This procedure is a mix of light mechanical and enzymatic treatment. For this purpose, a flat  
92 flexible silicon spatula and the enzyme zymolyase (Sigma-Aldrich, USA) were used. The  
93 zymolyase was used at the final concentrations of 5, 10, 20 and 25 U/ml. About 3 ml of each sample  
94 of macro-aggregates was transferred into a weighing vessel. The sample was gently and briefly  
95 (from seconds to 1 min) treated by crushing the aggregates in the weighing vessel with a flat silicon  
96 spatula to reduce their size. Then, for each sample four vials were prepared as follows: 200 µl of  
97 culture added to 200 µl of zymolyase solution (5, 10, 20, 25 U/ml). The samples were then observed  
98 under a microscope after the addition of 400 µl of Lugol's solution to fix the cells for 15, 30, 45 and  
99 60 min. Trials performed either with mechanical or enzymatic treatment were also carried out. To  
100 check the maintenance of cell vitality after the enzymatic treatment, before addition of the Lugol's

101 solution, a small quantity of each sample was inoculated on solidified (agar 1.5%) seawater and  
102 incubated at 25°C with a photoperiod of 18 h light/6 h dark for 20 days.

103 Bloom A was micromanipulated (MSM System 400, Singer Instrument Co Ltd, UK) in order to  
104 obtain purified species. Briefly, the culture was spread on one side of Petri plates containing  
105 solidified seawater (plus Walne and vitamins). Cells with different morphology were placed at  
106 defined positions on the plates (according to the MSM System instruction). The plates were then  
107 incubated at 25°C with a photoperiod of 18 h light/6 h dark. The two genera of microalgae that were  
108 isolated were grown in enriched (Walne plus vitamins) seawater for 14 days and cell counting was  
109 carried out.

110

## 111 **Results**

112 After one week of culturing, the three samples were composed of different genera of microalgae  
113 and associated populations of bacteria. During the development, the samples turned a green/brown  
114 colour and showed persistent macroscopic aggregation that impaired the normal study of the  
115 cultures.

116 The samples showed a natural evolution consisting of the appearance of some species and the  
117 disappearance of others until the establishment of stable mixed populations composed of four/six  
118 species for each bloom. According to the morphological features, bloom A was composed of  
119 *Extubocellulus* cf. *spinifer*, *Nitzschia* sp., *Navicula* sp. and *Nannochloropsis* sp. Bloom B was  
120 composed of *Pseudoanabaena* sp., *Cymbella* sp., *Nitzschia* sp. and *Navicula* sp. Bloom C was  
121 composed of *Pseudoanabaena* sp., *Nodularia* sp., *Nitzschia* sp., *Navicula* sp., *Cylindrotheca* sp. and  
122 *Nannochloropsis* sp. The population of bacteria normally associated with the natural population  
123 sample was more evident after the enzymatic treatment because the bacterial cells were freed from  
124 the microalgal matrix.

125 A reduction in size of the macro-aggregates was not observed with zymolyase treatment at  
126 concentrations of 5 and 10 U/ml; thus, neither concentration resulted in a disaggregation of different  
127 microalgal genera when observed under the microscope. At concentrations of 20 and 25 U/ml there  
128 was a visible reduction in size of the macro-aggregates. As an example, Figure 1 shows sample C  
129 treated with spatula and zymolyase at a concentration of 20 U/ml on glass slides. Compared with  
130 the microscopic features of the cultures observed before the mechanical plus enzymatic treatment,  
131 the best result was obtained using the enzyme concentration of 20 U/ml after 45 min, as shown in  
132 Figures 2–4 for the three blooms A, B and C observed under the microscope, respectively. The  
133 higher concentration of the enzyme did not produce any improvement compared to the  
134 concentration of 20 U/ml (Figure 5). Also, performing either mechanical or enzymatic treatment did  
135 not give satisfactory results, as shown for bloom B as an example (Figure 6). For all the  
136 experiments under the microscope the cells observed were intact; moreover, the three blooms  
137 succeeded in growing on solidified seawater plates, demonstrating that they were not negatively  
138 affected by the enzyme. Therefore, the treatment did not cause cell lysis. Figure 7 shows the growth  
139 curves of *Extubocellulus* cf. *spinifer* and *Nannochloropsis* sp. isolated from bloom A by  
140 micromanipulation.

141

## 142 **Discussion**

143 The proposed method facilitates the study of mixed microalgal populations, especially for  
144 laboratories that do not have expensive technologies available. The observation of intact algal cells  
145 is consistent with the results of other authors (Allard et al. 2002; Versteegh & Blokker 2004) due to  
146 the high resistance to degradation of most microalgal cell walls. Sander & Murthy (2009) reported  
147 that, depending on the species, the inner cell wall layer – often composed of cellulose – may be  
148 enclosed in layers of other substances such as mucilage, pectin, proteins, lipids and sulfonated  
149 polysaccharides that together can make the cell wall less susceptible to enzymes. This study

150 provided evidence of the maintenance of cell vitality after the enzymatic treatment and of the ability  
151 of the three bloom communities to grow on solidified seawater after treatment. The reduction in size  
152 of the aggregates could be determined by the zymolyase attacking the microalgal EPS. This study is  
153 an initial contribution that proposes an approach to study mixed algal populations. Different authors  
154 have reported on the diversity of cell walls in algae (Domozych et al. 2012; Popper et al. 2014) and  
155 of the EPS in various groups of microalgae (de Jesus Raposo et al. 2013), in cyanobacteria (Parikh  
156 & Madamwar 2006) and in diatoms (Chiovitti et al. 2003; Higgins et al. 2003). This approach takes  
157 into account the possible use of different enzymes according to the differences of cell walls and  
158 EPS of the cells in the mixed population. Moreover, for the optimal results of this proposed method,  
159 it is important to highlight the role of the light mechanical treatment coupled with the use of the  
160 enzymes. In fact, the brief mechanical treatment by crushing is needed to reduce the macro-  
161 aggregates (that still remain macroscopically visible) to enable the withdrawal of the sample as  
162 homogeneously as possible to be added to the enzyme. The treatment of the sample with the spatula  
163 was only effective for up to a minute as the size of the aggregates no longer varied. Also,  
164 performing the enzymatic treatment directly was not feasible because the aggregates were too large  
165 to be homogeneously taken up using normal pipette tips.

166 The proposed approach is useful for distinguishing the morphology of the different species  
167 present in a natural population, allowing for their identification by classical keys; furthermore, it  
168 allows a pure culture to be obtained from an inoculum of mixed species using a micromanipulator,  
169 and allows cell counting, particularly if the bloom does not consist of filamentous species.

170

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173 identification.

174 **Disclosure statement**

175 No potential conflict of interest was reported by the author.

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179 I-B/2.2/099.

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242 **Figure legend**

243 **Figure 1.** Marine microalgal mixed bloom (sample C) on glass slides (a) before and after (b) 15  
244 min, (c) 30 min, (d) 45 min and (e) 60 min of mechanical and zymolyase (20 U/ml) treatment.

245 **Figure 2.** Microscopic observation (300×) of the marine microalgal mixed bloom A (a) before and  
246 (b) after 45 min of mechanical and zymolyase (20 U/ml) treatment.

247 **Figure 3.** Microscopic observation (300×) of the marine microalgal mixed bloom B (a) before and  
248 (b) after 45 min of mechanical and zymolyase (20 U/ml) treatment.

249 **Figure 4.** Microscopic observation (300×) of the marine microalgal mixed bloom C (a) before and  
250 (b) after 45 min of mechanical and zymolyase (20 U/ml) treatment.

251 **Figure 5.** Marine microalgal mixed bloom (sample B): (a) on glass slide after 45 min of treatment  
252 with zymolyase (20 U/ml) and (b) observed under the microscope (300×); (c) on glass slide after 45  
253 min of treatment with zymolyase (25 U/ml) and (d) observed under the microscope (300×).

254 **Figure 6.** Marine microalgal mixed bloom (sample B) treated either by (a) mechanical treatment or  
255 (b) by enzymatic treatment (20 U/ml) after 45 min.

256 **Figure 7.** Growth curves and microscopic images (300×) of (a) *Extubocellulus cf. spinifer* and (b)  
257 *Nannochloropsis* sp. isolated from bloom A by micromanipulation.

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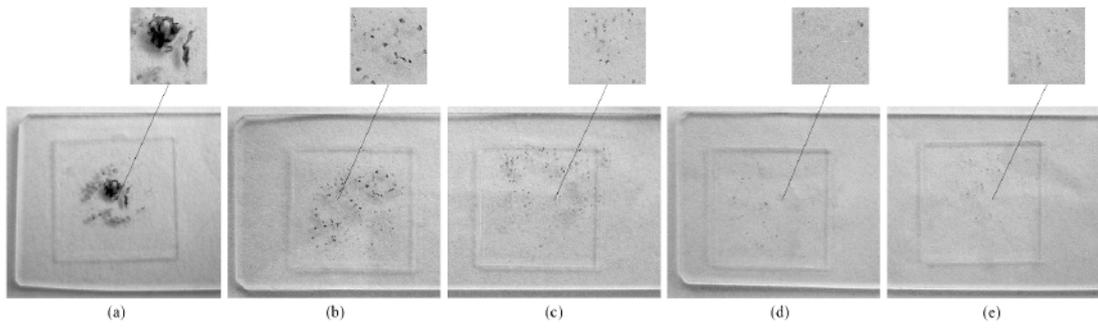
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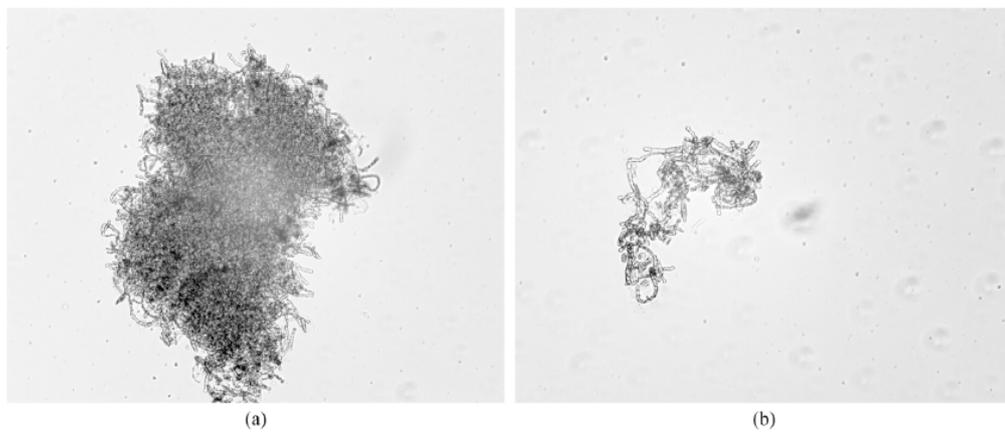


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**Fig. 1**

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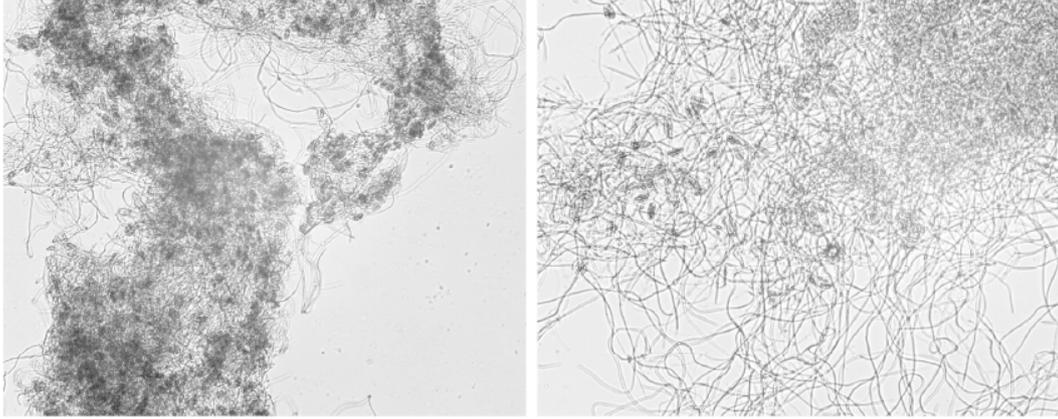
**Fig. 2**

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(a)

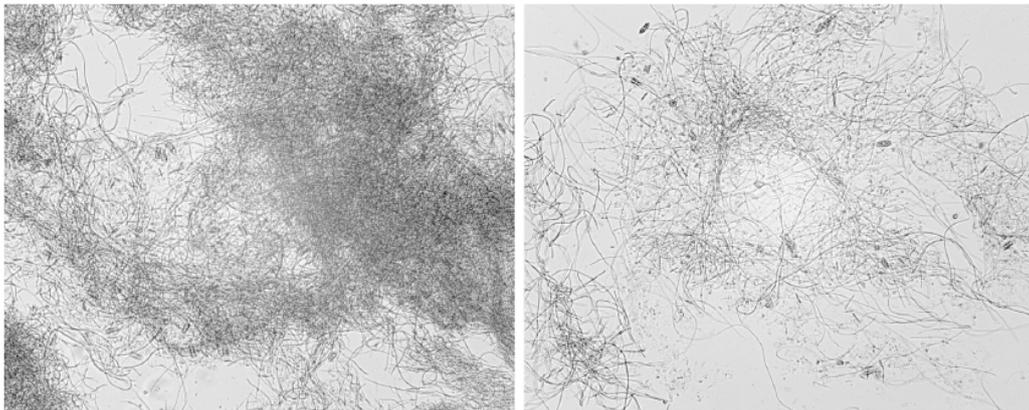
(b)

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**Fig. 3**

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(a)

(b)

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**Fig. 4**

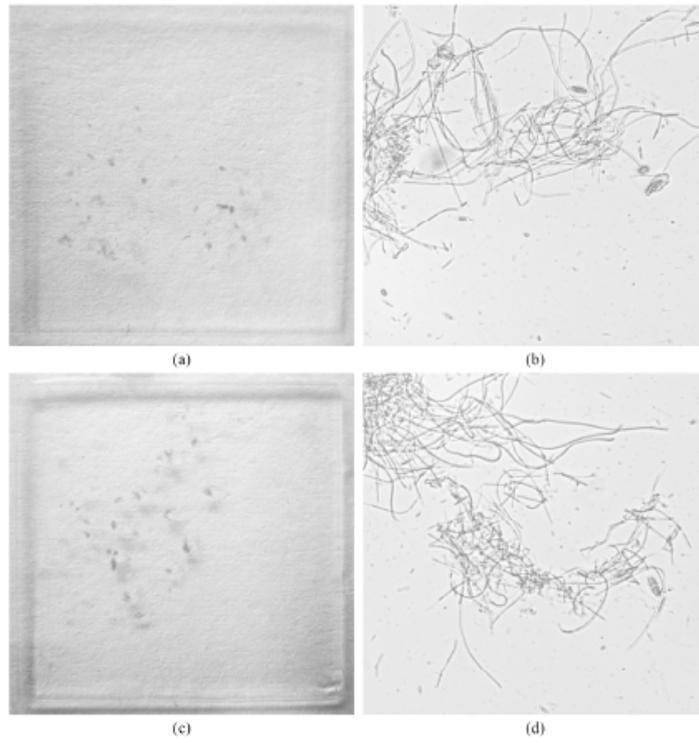
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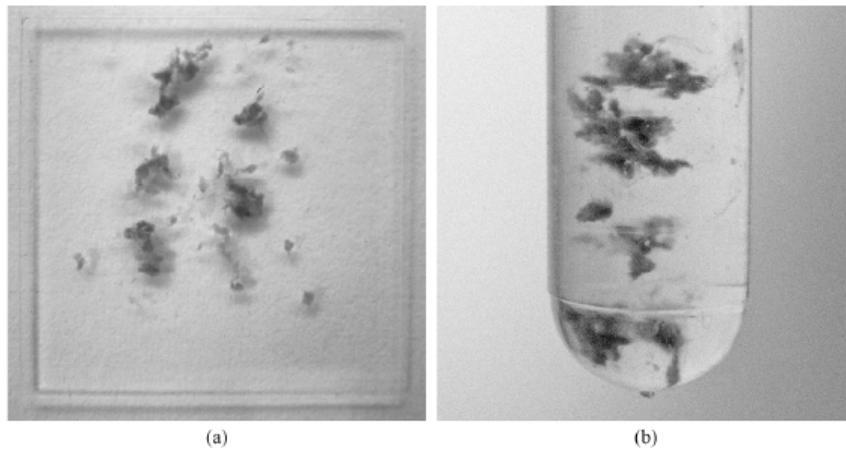


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**Fig. 5**

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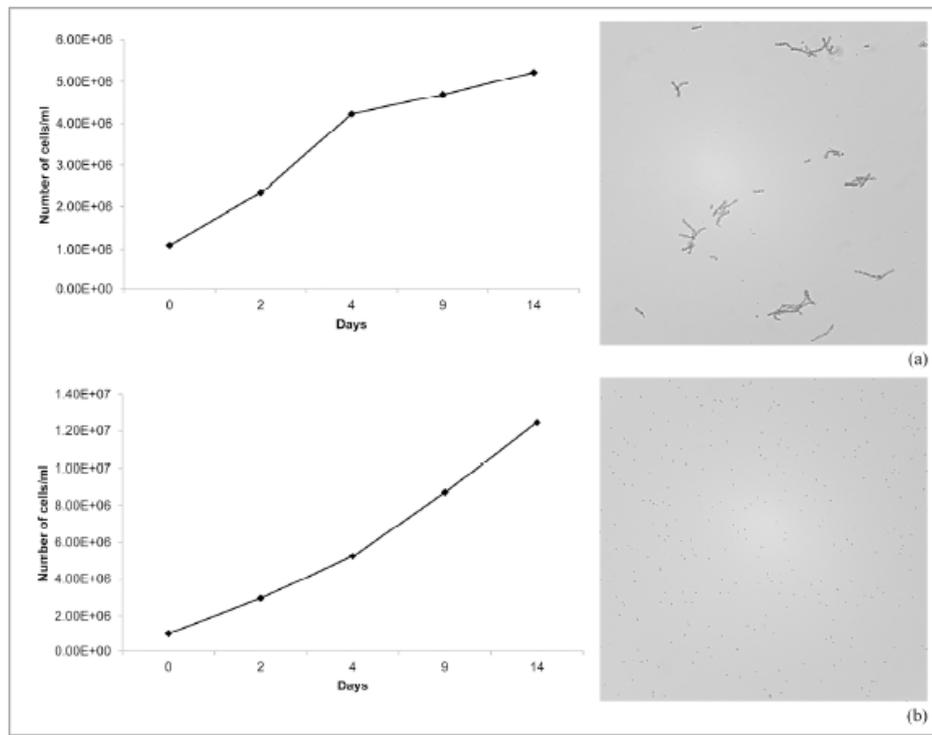
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**Fig. 6**

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**Fig. 7**