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## Biotechnological properties and antifungal activity of lactic acid bacteria isolated from two marine algae *Ulva lactuca* and *Sargassum muticum* collected from the Moroccan coast of Sidi Bouzid-EI Jadida

--Manuscript Draft--

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| <b>Abstract:</b>                                     | <p>Isolation of new strains of lactic acid bacteria (LAB) from extreme environments is currently in great demand given their biotechnological properties and their potential integration into food biotransformation and valorization processes. Thus, the purpose of this work was to isolate LAB strains from two marine algae (<i>Ulva lactuca</i> and <i>Sargassum muticum</i>) collected from the coast of Sidi Bouzid-EI Jadida (Morocco). A total of 15 potential LAB strains were isolated on MRS agar on the two marine algae species. Microscopic and phenotypic identification of LAB isolates showed that all strains were Gram-positive, cocci-shaped and showed negative catalase reaction. Biochemical tests showed that all LAB isolates were homofermentative and 10 LAB strains (66.6%) showed lipolytic and/or proteolytic enzymatic activities. Out of the 15 LAB isolates, 8 strains exhibited a considerable antifungal activity against fungal strains of <i>Aspergillus flavus</i> and <i>A. niger</i> ranging from 60 to 90%. The ability of LAB isolates to grow on culture media formulated by the two algae powders showed a significant acidifying capacity and a decrease in pH from an initial value of 7.04 to 2.63 after growth for 10 days. These results suggest the potential use of LAB isolates in fermentation and biotechnological processes for the valorization of marine algae into value-added products for various applications.</p> |                         |
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1  
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4 **collected from the Moroccan coast of Sidi Bouzid-El Jadida**

5  
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12  
13 **Abstract**

14 Isolation of new strains of lactic acid bacteria (LAB) from extreme environments is currently  
15 in great demand given their biotechnological properties and their potential integration into food  
16 biotransformation and valorization processes. Thus, the purpose of this work was to isolate  
17 LAB strains from two marine algae (*Ulva lactuca* and *Sargassum muticum*) collected from the  
18 coast of *Sidi Bouzid-El Jadida* (Morocco). A total of 15 potential LAB strains were isolated on  
19 MRS agar on the two marine algae species. Microscopic and phenotypic identification of LAB  
20 isolates showed that all strains were Gram-positive, cocci-shaped and showed negative catalase  
21 reaction. Biochemical tests showed that all LAB isolates were homofermentative and 10 LAB  
22 strains (66.6%) showed lipolytic and/or proteolytic enzymatic activities. Out of the 15 LAB  
23 isolates, 8 strains exhibited a considerable antifungal activity against fungal strains of  
24 *Aspergillus flavus* and *A. niger* ranging from 60 to 90%. The ability of LAB isolates to grow  
25 on culture media formulated by the two algae powders showed a significant acidifying capacity  
26 and a decrease in pH from an initial value of 7.04 to 2.63 after growth for 10 days. These results  
27 suggest the potential use of LAB isolates in fermentation and biotechnological processes for  
28 the valorization of marine algae into value-added products for various applications.

29 **Keywords:** Marine algae, *Ulva lactuca*, *Sargassum muticum*, Lactic acid bacteria, Antifungal,  
30 Fermentation, Valorisation, Biotechnological process.

31  
32 **Introduction**

33 Global revolution in food production, social and environmental sustainability, health, nutrition  
34 and economic development has increased attention to marine resources, including the  
35 production and use of algae. Compared to terrestrial plants, marine algae can grow faster and  
36 year-round, resulting in multiple harvests per year and higher yields per unit area. Marine algae  
37 are considered a rich resource of various bioactive compounds, such as antimicrobial,  
38 antioxidant, antiproliferative and anticancer substances (Hmani et al. 2021). In addition, marine  
39 algae production has many environmental advantages over land-based agriculture, such as not

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40 needing fertilizer, pesticides or fresh water. However, traditional techniques used to extract  
41 bioactive compounds from marine algae such as acid hydrolysis using high temperature or using  
42 chemical solvent extraction, are often costly and produce large amounts of wastes, some of  
43 which could be complex and may be toxic or have negative impact on the environment. On the  
44 other hand, heat-sensitive compounds, as bioactive compounds (phenolic compounds,  
45 carotenoids, etc.) may be degraded during algae extraction process. As a result, green  
46 technologies such as fermentation and enzyme treatment have emerged as alternatives to  
47 traditional extraction techniques because they can obtain the desired compounds without  
48 producing toxic residues or using solvents that may alter the bioactive compounds activity  
49 (Pérez-Alva et al. 2022).

50 Fermentation of marine algae with LAB strains has recently been reported in the literature, and  
51 the use of LAB strains for the fermentation of algal by-products to produce lactic acid is  
52 considered a promising approach. Furthermore, marine algae are considered as important  
53 natural resources of bioactive substances such as phenolic compounds, carotenoids, vitamins,  
54 proteins, lipids and polysaccharides (Sumardianto Riyadi et al. 2021). Moreover, marine algae  
55 are renewable substrates rich in fermentable sugars, with green and red algae (macro-algae)  
56 generally containing 30-50% complex sugars that are fermentable after pretreatment.

57 Despite the abundance of scientific information in the literature on LAB isolated from terrestrial  
58 matrices especially from cereal products, vegetables, plants, and fruits, etc., few studies are  
59 available on LAB strains isolated from marine origin and their use for marine algae  
60 fermentation and transformation. Indeed, lactic acid production from algae was described for  
61 the first time in Japan but is still poorly studied. Currently, marine algae are mainly used in food  
62 and feed, for the production of algae-based fertilizers, biofuels, and bioplastics. However, the  
63 use of algae as culture medium for lactic acid production by homofermentative LAB is less  
64 studied compared to conventional substrates, since homofermentative LAB produce 2 times  
65 more lactic acid than the heterofermentative LAB (Hwang et al. 2012, Sudhakar and Dharani  
66 2022). According to recent studies, many algal species could be used as good substrates for  
67 LAB fermentation, and may open the possibility of obtaining many add value products to be  
68 incorporated as new ingredients in human diets, animal feed or to be used as new organic soil  
69 fertilizers, etc. (Uchida and Miyoshi 2013).

70 Morocco, a North African and Mediterranean country, with a total of 3500 km of coastline, is  
71 considered as a country with the longest coastline on the African continent. Moroccan coasts  
72 are particularly rich in marine algae and constitute a reserve of species with economic, social  
73 and ecological potential. Previous studies have reported the isolation of several LAB species  
74 from different matrices as example of raw camel milk (Khedid et al. 2009), poultry waste  
75 manure (El-Jalil et al. 2008) and sardines (Ndaw et al. 2008). Thus, this study aimed to isolate,  
76 for the first time, new strains of LAB from two abundant marine algae *Ulva lactuca* and  
77 *Sargassum muticum* collected from the Moroccan coast of *Sidi Bouzid* at El Jadida city, as well  
78 as, to evaluate their multifunctional properties (biotechnological and antifungal) for their  
79 potential application in various biotechnological processes.

80

## 81 **Material and Methods**

82

### 83 **Algae sampling and preparation**

84 Two species of abundant marine algae (*Ulva lactuca* and *Sargassum muticum*) were collected  
85 on the coast of *Sidi Bouzid* in El Jadida city (Morocco) during the period April-June 2022. After

**Brahimi et al. Biotechnological properties and antifungal activity of lactic acid bacteria isolated from two marine algae *Ulva lactuca* and *Sargassum muticum* collected from the Moroccan coast of Sidi Bouzid–El Jadida. Euro–Mediterranean Journal for Environmental Integration.**

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86 washing with tap water and distilled water for the removal of salts, algae samples were dried at  
87 room temperature (25°C), then ground with a Blender laboratory, and finally the ground  
88 material was then sieved to obtain a fine algae powder and stored at 4°C until analysis.

89

### 90 **Isolation and characterization of LAB**

91 One (1 g) of each algae powder sample was mixed with 10 mL of MRS broth (De Man, Rogosa  
92 and Sharpe, Biokar, France) and incubated at 30°C for 24h under aerobic conditions. LAB  
93 strains isolation was carried-out on MRS agar supplemented with bromocresol purple (0.002  
94 %) as a color indicator and yellow colonies were selected and purified (Shobharani et al. 2012).  
95 Potential LAB isolates were examined macroscopically based on the morphology of colonies,  
96 microscope observation and catalase reaction (Prastujati et al. 2022). Gram-positive LAB  
97 isolates tested as catalase negative were stored on MRS slant at 4°C for further biochemical  
98 characterization.

99

### 100 **Growth at different temperature values**

101 LAB isolates were inoculated in sterile MRS broth tubes and incubated at different temperature  
102 values (30°C, 37°C and 45°C) for 48 h to distinguish mesophilic LAB strains from thermophilic  
103 LAB strains (Belkacem et al. 2009).

104

### 105 **CO<sub>2</sub> and exopolysaccharides (EPS) production**

106 The homo- or heterofermentative properties of LAB isolates were studied based on their ability,  
107 or not, to produce CO<sub>2</sub> on MRS semi-agar enriched with sugar (5% glucose) after incubation at  
108 30°C for 48 h under aerobiosis. For EPS production, LAB isolates were streaked onto MRS  
109 agar medium supplemented with 5% of glucose and incubated aerobically at 30°C for 48 h.  
110 EPS production was considered positive if colonies showed mucoid or viscous shapes on the  
111 agar medium (Khalil et al. 2018).

112

### 113 **Lipolytic and proteolytic activity**

114 Proteolytic activity of the LAB isolates was assessed by depositing 20 µL of LAB cultures on  
115 sterile Whatman paper disks, previously placed on the surface of MRS agar supplemented with  
116 10% skimmed milk. Proteolytic activity was evaluated as a clear zone around the discs after  
117 incubation at 30° C for 24 h under aerobiosis, while lipolytic activity was tested by inoculating  
118 LAB isolates on agar spot (MRS supplemented with 1% of tributyrin). After incubation at 30°  
119 C for 72 h under aerobic conditions, positive lipolytic activity was elucidated as an opaque zone  
120 around the disc due to the formation of fatty acids and esters released by calcium.

121

### 122 **Antifungal activity**

123 The antifungal activity of LAB isolates was tested against the growth of fungal strains of *A.*  
124 *flavus* and *A. niger* (fungal collection of BIOMARE Laboratory, UCD University, El Jadida).  
125 Mycelium growth inhibition by the cell-free supernatants (CFS) of LAB strains was determined  
126 on Dichloran Rose-Bengal Chloramphenicol (DRBC) Agar (Biokar, France). Plates containing  
127 DRBC medium, added at 10% (v/v) with sterile CFS of each LAB strain, were inoculated in  
128 the center with disc (5 mm) of *A. flavus* and *A. niger*. After aerobic incubation at 25°C for 5  
129 days, the diameters of fungal colonies were measured. A negative control was used by adding  
130 MRS broth to DRBC medium (Wang et al. 2002).

131

*Brahimi et al. Biotechnological properties and antifungal activity of lactic acid bacteria isolated from two marine algae Ulva lactuca and Sargassum muticum collected from the Moroccan coast of Sidi Bouzid–El Jadida. Euro–Mediterranean Journal for Environmental Integration.*

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## 132 LAB application for algae fermentation

133

### 134 Screening of LAB growth on algae media

135 In order to test the growth of LAB isolates of algae-based media, two culture media were  
136 separately prepared using all MRS components, except glucose that was replaced with algae  
137 powder (*S. muticum* or *U. lactuca*) as carbon sources. Since LAB strains are demanding in terms  
138 of nutritional substances, the first objective of this study was to check if LAB isolates are  
139 capable of using algae-carbohydrate. For this, LAB strains were inoculated on the algae media  
140 (*S. muticum* culture medium or *U. lactuca* culture medium) and incubated at 30°C for 48 h  
141 under aerobic conditions, as previously reported by Shobharani et al. (2012).

142

### 143 Algae pre-treatment and fermentation assay

144 Algae fermentation assays with LAB strains were carried out by preparing a hydrolysate from  
145 the two studied algae species. Thus, 2 g of each algae powder were mixed with 100 mL of  
146 distilled water and added to 1% of H<sub>2</sub>SO<sub>4</sub> (Sudhakar and Dharani 2022). The mixture was  
147 sterilized at 121°C for 20 min to obtain a juice extract from each algae species. The pre-treated  
148 algal extracts potentially rich in polysaccharides were then filtered and supplemented with the  
149 following nutrients (g/L): yeast extract (5.5), peptone (12.5), K<sub>2</sub>HPO<sub>4</sub> (0.25), KH<sub>2</sub>PO<sub>4</sub> (0.25),  
150 CH<sub>3</sub>COONa (10), MgSO<sub>4</sub> (0.1), MnSO<sub>4</sub> (0.05 mg), FeSO<sub>4</sub> (0.05 mg) and the pH was adjusted  
151 to 7. Two algae extract media (AEM) were prepared and sterilized at 121 °C for 15 min. Later,  
152 LAB strains isolated from each algal species were grown on the AEM medium prepared by the  
153 respective algae powder. Indeed, 50 mL each prepared AEM were inoculated with a preculture  
154 of each LAB strain at 10% (v/v), and incubated at 30°C for 10 days under aerobiosis. The pH  
155 evolution, as an indicator of LAB growth was followed during the fermentation assay of each  
156 AEM. This step is required for the selection of the most growing and acidifying isolates on  
157 AEM. Performing strains will be selected to be used as starters for bulk algae fermentation.

158

### 159 Statistical analysis

160 All experiments in this study were performed in triplicate (n = 3). The Student t test was used  
161 to confirm the significant difference between the two means of inhibition activity of LAB  
162 isolated from the algae studied against the two fungal strains tested. The difference was  
163 significant at p<0.05. The digital tool to perform this statistical analysis is the Excel 2013  
164 software. Concerning the analysis by the principal component analysis (PCA) method and LAB  
165 strains clustering, we used the free software Past 4.

166

## 167 Results and discussion

168

### 169 LAB isolation

170 This first step consisted of a phenotypic identification of cultivable LAB strains isolated from  
171 the two algae powders. Purification of isolates on the MRS culture medium resulted in fifteen  
172 (15) pure strains. It should be highlighted that all studied strains have cocci forms and there  
173 were no lactobacilli-form among the isolates. Furthermore, all strains were Gram positive and  
174 catalase negative as main characteristics. These traits suggest their probable membership in  
175 lactic acid bacteria group.

176

### 177 Biochemical properties of LAB isolates

**Brahimi et al.** Biotechnological properties and antifungal activity of lactic acid bacteria isolated from two marine algae *Ulva lactuca* and *Sargassum muticum* collected from the Moroccan coast of Sidi Bouzid–El Jadida. Euro–Mediterranean Journal for Environmental Integration.

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178 Results of the growth at different temperature values showed that out of the 15 strains tested, 8  
179 strains (S1, S6, S3, S5, S7, U2, U3, U5) were able to grow at 45°C. LAB growth at this  
180 temperature is manifested by the presence of a bacterial cloud (ring) at the bottom of the tubes  
181 when compared to the control. According to Boudjema et al. (2009), LAB are known for their  
182 good lactic acid production (9.06 g/L) at temperature values of 42-45°C, compared to other  
183 temperature growth values. On the other hand, all tested LAB isolates belong to the  
184 homofermenters group. Given that isolated strains are mesophilic and others are thermophilic,  
185 we can suggest that mesophilic bacteria could belong to the *Streptobacterium* group, while  
186 thermophilic bacteria belong to *Thermobacterium* group (Ruiz Rodríguez et al. 2019).

187

### 188 **Biotechnological properties of LAB**

189 Studying the biotechnological properties of LAB is essential to characterize the isolates for their  
190 potential use in future laboratory and large-scale pilot applications. Properties of LAB isolated  
191 from the two studied marine algae species are summarized in table 1.

192 Regarding EPS production, all LAB strains (except S5 and S10) were positive. These results  
193 showed that algae powders are rich in LAB, from which EPS-producing strains can be isolated.  
194 EPS production by LAB strains varies from strain to strain and is considered as one of the most  
195 sought after and desired biotechnological properties since this property plays a critical role on  
196 the texture of fermented products. Some LAB strains, belonging to genera *Leuconostoc*,  
197 *Weissella* and *Pediococcus*, are able to synthesize and excrete EPS from sucrose by  
198 extracellular glucansucrases or fructansucrases (İspirli et al. 2020). It was reported that EPS  
199 production by LAB is considered as one of the most desired biotechnological properties since  
200 EPS play a crucial role on the texture of fermented matrices (EL Boujamaai et al. 2023). Our  
201 results are similar to those that already reported the production of EPS by LAB strains such as  
202 *Enterococcus faecium*, *Limosilactobacillus fermentum*, *Lacticaseibacillus casei*, *Leuconostoc*  
203 *mesenteroides* and *Streptococcus thermophilus*, etc. (Zannini et al. 2016; Lynch et al. 2018).  
204 Differences between LAB strains in EPS production suggest that EPS amounts is a strain-  
205 dependent characteristic, influenced by the bacterial strain and growth phase, including the  
206 environmental conditions of the culture medium. These conditions include carbon source,  
207 nitrogen and physicochemical conditions for bacterial growth, especially temperature, pH,  
208 oxygen levels, etc. (Khalil et al. 2018).

209 Regarding enzyme production by LAB isolates, the lipolytic activity showed that out of 15 LAB  
210 strains, 11 isolates (73.3%) have significant lipolytic activity in the presence of 1% tributyrin.  
211 It is well known that LAB are generally considered to be weakly lipolytic, as compared to other  
212 groups of microorganisms. It should be highlighted that algae contain varied forms of lipids  
213 owing to their extreme habitat diversity, especially n-6 and n-3 polyunsaturated fatty acids  
214 (PUFAs), glycolipids, and phospholipids and sterols (Kumari et al. 2013). These lipids are  
215 known with high nutritional value and with important bioactive properties (Domingues and  
216 Calado 2022). Thus, the presence of a significant lipolytic activity in LAB isolated from marine  
217 algae could be explained by their adaptation to their natural substrates (marine algae)  
218 characterized by a richness in lipids, where the lipolytic activity of such LAB could play an  
219 important role during algae fermentation by LAB isolates. In comparison, previous studies  
220 isolated 29 lipase-producing strains from over 100 different LAB representing the genera of  
221 *Lactobacillus*, *Lactococcus*, *Leuconostoc*, *Pediococcus* and *Streptococcus* (Meyers et al.,  
222 1996). Recently, Aziz et al. (2021, 2022) reported the ability of *L. plantarum* 12-3 to convert

**Brahimi et al.** Biotechnological properties and antifungal activity of lactic acid bacteria isolated from two marine algae *Ulva lactuca* and *Sargassum muticum* collected from the Moroccan coast of Sidi Bouzid–El Jadida. Euro–Mediterranean Journal for Environmental Integration.

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223 linoleic acid at different concentrations to different health-beneficial fatty acid and other  
224 metabolites.

225  
226  
227

**Table 1** : Metabolic properties and phenotypic characteristics of the isolated LAB strains

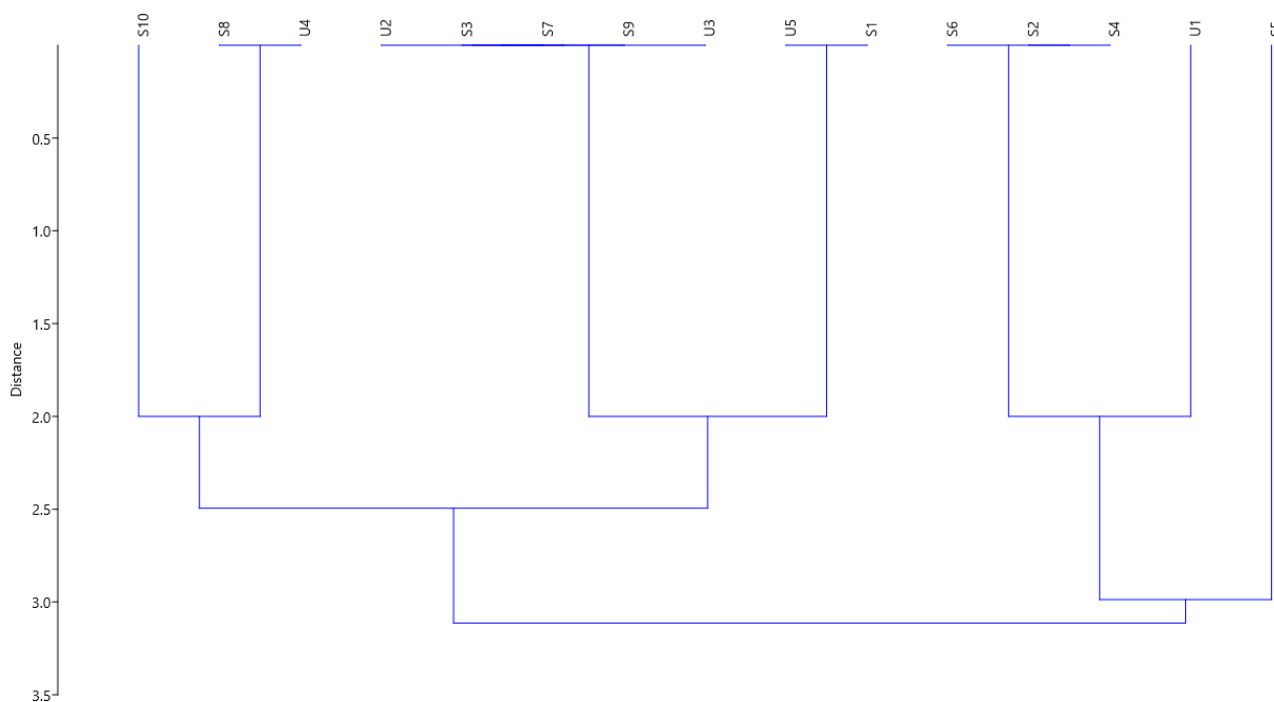
| Studied algae     | LAB Strains code | Growth temperature (°C) |     |     | Metabolic properties |     |      |     |
|-------------------|------------------|-------------------------|-----|-----|----------------------|-----|------|-----|
|                   |                  | 30                      | 37  | 44  | EPS                  | Lip | Prot | Hom |
| <i>S. muticum</i> | S1               | (+)                     | (+) | (+) | (+)                  | (+) | -    | (+) |
|                   | S2               | (+)                     | (+) | -   | (+)                  | -   | -    | (+) |
|                   | S3               | (+)                     | (+) | (+) | (+)                  | (+) | (+)  | (+) |
|                   | S4               | (+)                     | (+) | -   | (+)                  | -   | -    | (+) |
|                   | S5               | (+)                     | (+) | (+) | -                    | -   | -    | (+) |
|                   | S6               | (+)                     | (+) | -   | (+)                  | -   | -    | (+) |
|                   | S7               | (+)                     | (+) | (+) | (+)                  | (+) | (+)  | (+) |
|                   | S8               | (+)                     | (+) | -   | (+)                  | (+) | (+)  | (+) |
|                   | S9               | (+)                     | (+) | (+) | (+)                  | (+) | (+)  | (+) |
|                   | S10              | (+)                     | (+) | -   | -                    | (+) | (+)  | (+) |
| <i>U. lactuca</i> | U1               | (+)                     | (+) | -   | (+)                  | (+) | -    | (+) |
|                   | U2               | (+)                     | (+) | (+) | (+)                  | (+) | (+)  | (+) |
|                   | U3               | (+)                     | (+) | (+) | (+)                  | (+) | (+)  | (+) |
|                   | U4               | (+)                     | (+) | -   | (+)                  | (+) | (+)  | (+) |
|                   | U5               | (+)                     | (+) | (+) | (+)                  | (+) | -    | (+) |

228 *EPS: Exopolysaccharides; Lip : lipolytic activity; Prot : Proteolytic activity ; Hom :*  
229 *Homolactic fermentation, (+): positive reaction; -: negative reaction*

**Brahimi et al.** Biotechnological properties and antifungal activity of lactic acid bacteria isolated from two marine algae *Ulva lactuca* and *Sargassum muticum* collected from the Moroccan coast of Sidi Bouzid–El Jadida. *Euro–Mediterranean Journal for Environmental Integration*.

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230  
 231 Results of the proteolytic activity showed that out of 15 isolates, 8 LAB (5.33%) exhibited a  
 232 positive activity. A bacterial strain is called proteolytic if it has a lysis zone of diameter between  
 233 5 and 15 mm. It therefore appears that the strains S3, S7, S8, S9, S10, U2, U3, and U4 are  
 234 strongly proteolytic. LAB strains U1 and U5 revealed a moderate proteolytic activity, since the  
 235 diameter of the proteolysis zones were between 9 mm and 12 mm, and the others (S1, S2, S4,  
 236 S5, S6, U1 and U5) are not proteolytic since the clear zone does not exceed 3 mm. Marine algae  
 237 are considered a viable source of protein and some species are known to contain protein levels  
 238 similar to those of traditional protein sources (meat, egg, soybean, and milk), which would  
 239 constitute alternative protein sources required to fulfil the consumers demand and to meet  
 240 predicted global protein requirements (Bleakley and Hayes 2017). Proteolytic activity of LAB  
 241 is a very important property because it is necessary for LAB growth in fermented matrices and  
 242 improving the organoleptic properties (texture and flavor) of these products (Zareie et al. 2023).  
 243 The analysis of distances between the characteristics illustrated in table 1 relating to LAB  
 244 strains isolated from the two algae species (*S. muticum* and *U. lactuca*) allowed the  
 245 neighborhood grouping represented in figure 1.  
 246



247  
 248 **Figure 1** : Neighborhood grouping of LAB strains isolated from the studied algal species.

249  
 250  
 251 From this illustration, it can be seen that the bacteria are classified into two groups, namely a  
 252 small group combining strains U1, S2, S4, S5 and S6 and a large group formed of strains U2,  
 253 U3, U4, U5, S1, S3, S7, S8, S9 and S10. In the first group, the strains S2, S4 and S6 are brought  
 254 closer to the strain U1 and the strain S5 forms a singleton which remains linked to the others  
 255 by the base of the classification branch. The large group according to data shown in figure 1 is



**Brahimi et al.** Biotechnological properties and antifungal activity of lactic acid bacteria isolated from two marine algae *Ulva lactuca* and *Sargassum muticum* collected from the Moroccan coast of Sidi Bouzid–El Jadida. Euro–Mediterranean Journal for Environmental Integration.

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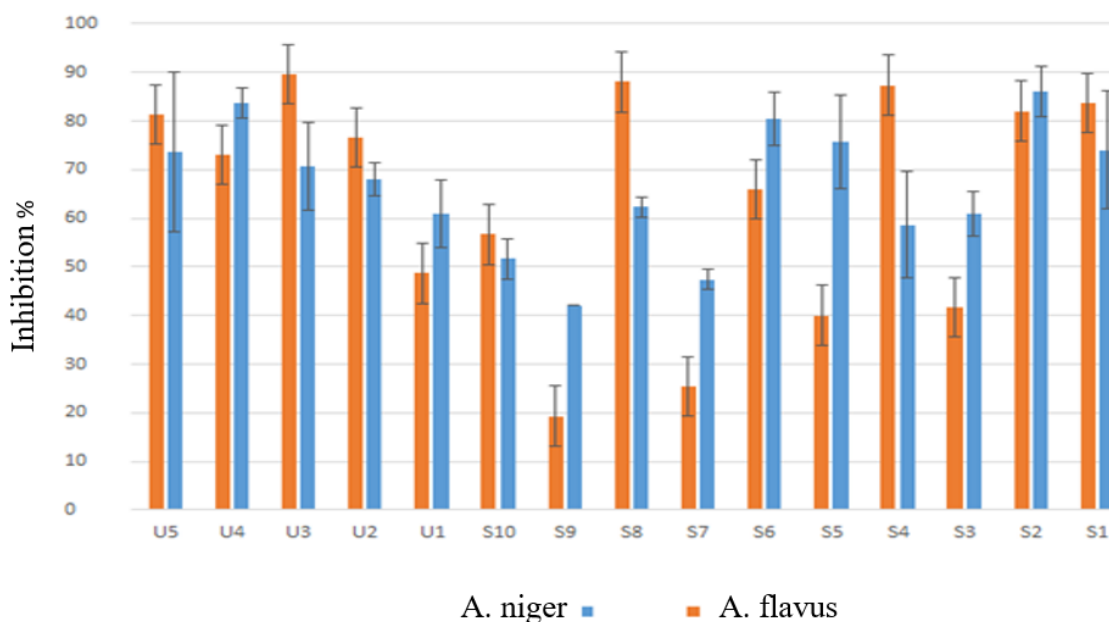
256 formed by the strains U4, S8 and S10 which form a small group and the strains S1, U2, U3, U5,  
 257 S7 and S9. This classification reflects a metabolic and taxonomic link between the isolated LAB  
 258 strains. The results illustrated in Figure 1 show that the isolated strains of the two algae share  
 259 certain genetic characteristics that allow them to live and survive in the small niches formed by  
 260 the two algae separately.

261

262 **Antifungal activity of LAB**

263 Currently, food preservation is based on the use of chemicals that can have adverse  
 264 consequences on consumer health (Leyva Salas et al. 2017). To this end, scientists are always  
 265 looking for natural compounds and beneficial microorganisms, such as new LAB strains or  
 266 their metabolites, as alternatives to avoid chemical additives use during food processing.  
 267 Results on the antifungal activity of LAB against the two fungal species of *A. niger* and *A.*  
 268 *flavus* showed that out of 15 total LAB strains, 8 strains (S1, S2, S6, S8, U2, U3, U4 and U5)  
 269 showed considerable antifungal activity (> 60%) against *A. flavus* and/or *A. niger* (Figure 2).  
 270 while the other LAB strains showed lower to moderate antifungal activity. Moreover, all LAB  
 271 isolates are acidifying, suggesting that their antifungal activity is due to the acidification in  
 272 addition to the potential production of other molecules active against the tested fungal species.  
 273 Similar results were reported by Aderiye (2010), who reported that 68% of LAB isolated from  
 274 Nigerian fermented foods had antifungal activity against *A. niger* and *A. flavus*. These results  
 275 are in agreement with those already reported by Muhialdin et al. (2018), Abouloifa et al. (2020),  
 276 and Ben Salah-Abbès et al., 2021) on the ability of LAB to inhibit the growth of various fungal  
 277 species, and to reduce mycotoxin (Belgacem et al. 2019; Badji et al., 2023). Indeed, LAB strains  
 278 are able to produce multiple antimicrobial compounds, including organic acids (lactic, acetic,  
 279 citric, formic and succinic acids, among others), bacteriocins, phenyllactic acid, hydroxy-fatty  
 280 acids, proteinaceous compounds, reuterin, and many other compounds with antimicrobial  
 281 effects (Bartkiene et al. 2019; Matevosyan et al. 2020).

282



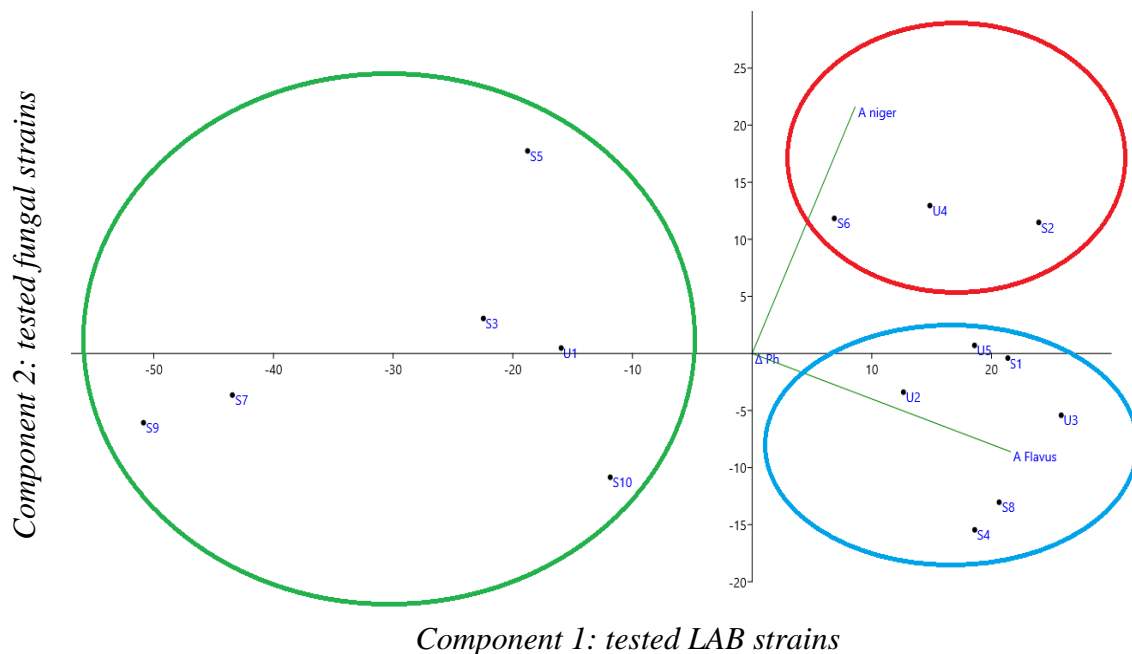
283

**Brahimi et al.** Biotechnological properties and antifungal activity of lactic acid bacteria isolated from two marine algae *Ulva lactuca* and *Sargassum muticum* collected from the Moroccan coast of Sidi Bouzid–El Jadida. Euro–Mediterranean Journal for Environmental Integration.

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284 **Figure 2** : Antifungal activity expressed as % inhibition of *A. niger* and *A. flavus* fungal  
285 growth.  
286

287 Dimensional analysis (Figure 3) performed by the principal component analysis (PCA)  
288 technique illustrates an antifungal activities distribution done by LAB strains (S1 to S10 and  
289 U1 to S5) isolated from *S. muticum* and *Ulva lactuca* algae as well than those of fungal strains  
290 tested (*A. niger* and *A. flavus*). According to figure 4, the LAB strains contribute in negative  
291 and positive parts of component 1 while the fungal strains contribute in the positive part (*A.*  
292 *niger*) and the negative part (*A. flavus*) of component 2. pH variation ( $\Delta$  pH) contributes to both  
293 components (1 and 2) at level 0. Figure 4 also shows that the strain *A. niger* is correlated  
294 (sensitive) to strains U4, S2 and S6 and the strain *A. flavus* is correlated (sensitive to strains U2,  
295 U3, U5, S1, S4 and S8). The other strains form a third group not correlated with the fungal  
296 strains tested.  
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304  
305 **Figure 3:** Dimensional analysis of antifungal activities of LAB strains (S1 to S10 and U1 to  
306 U5) isolated from *S. muticum* and *U. lactuca*.  
307  
308

309 **Application of LAB for algae fermentation**

310  
311 **LAB growth on algae media**  
312

**Brahimi et al.** Biotechnological properties and antifungal activity of lactic acid bacteria isolated from two marine algae *Ulva lactuca* and *Sargassum muticum* collected from the Moroccan coast of Sidi Bouzid–El Jadida. Euro–Mediterranean Journal for Environmental Integration.

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313 The growth of LAB isolates on modified MRS agar media was assessed by replacing the  
314 amount of glucose in the conventional MRS medium with an equivalent amount of marine algae  
315 powder of *S. muticum* or *U. lactuca*. Results showed a good growth of all LAB isolates on the  
316 AEM (both agar and broth).

317

### 318 pH monitoring during fermentation

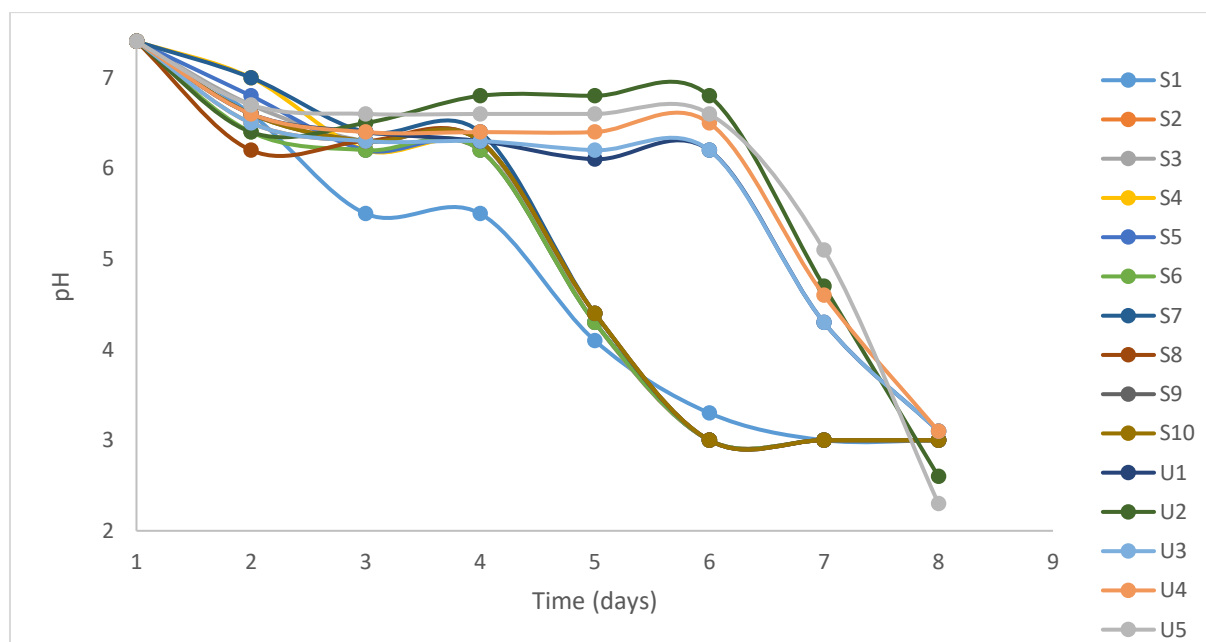
319 LAB isolates were firstly tested on algae autolysate and strong bacterial growth was observed  
320 at the bottom of the tubes, which allowed to conclude about the ability of LAB strains to growth  
321 on the algae based medium and to ferment algal components, especially algae polysaccharides.  
322 Fermentation assays were performed by pH measurement. Results of this experiment are  
323 reported in figure 4. As shown, all LAB strains were able to acidify AEM (prepared using *U.*  
324 *lactuca* or *S. muticum* powder) by lowering pH from an initial value of 7.4 to a final pH value  
325 of 3 for LAB strains S1, S6 and S10 isolated from *S. muticum* strains, and to a final pH value  
326 2.3 and 2.6, respectively, for LAB strains U5 and U2 isolated from *U. lactuca*. It should be  
327 highlighted that all reached pH values are close or below to a pH value of 3, which be explained  
328 by the very good growth of isolated strains on the AEM and the high production of organic  
329 acids, mainly lactic acid, by LAB isolates during the fermentation assay.

330

331 As shown in figure 4, it can be noted that the pH drop follows a bimodal phenomenon (a slow  
332 phase and a fast phase). The LAB strains (U1 to U5) isolated from *U. lactuca* present a first  
333 slow phase for 6 days then a linear fast phase. The linear model identified in the strains isolated  
334 from the *U. lactuca* presents a long organic acid production metabolism. The drop in pH values  
335 only occurred after the 6<sup>th</sup> day of growth. This rapid drop resulted in low record pH values,  
336 especially for the LAB strains U2 and U5 (*U. lactuca*) that decreased the pH at the end of  
337 bacterial growth until very low values of 2.6 and 2.3, respectively.

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**Figure 4.** pH evolution of AEM by LAB strains isolated from the studied algal species.

**Brahimi et al.** Biotechnological properties and antifungal activity of lactic acid bacteria isolated from two marine algae *Ulva lactuca* and *Sargassum muticum* collected from the Moroccan coast of Sidi Bouzid–El Jadida. *Euro-Mediterranean Journal for Environmental Integration*.

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343  
344

345 Whereas the LAB strains isolated from *S. muticum* present a first slow phase which lasts only  
346 4 days followed by a rapid logarithmic phase. These strains exhibit a remarkable adaptation  
347 to the massive production of organic acids by fermentable sugars bioconversion (Straathof  
348 2023). Indeed, the strain S1 (*S. muticum*) resulted in a drastic extreme decrease from an initial  
349 pH value of 7.4 to final pH values of 4.1 and 3.1 after 5 and 6 days of growth, respectively.  
350 This pattern has been observed with other LAB strains isolated from *S. muticum*, especially the  
351 strains S6 and S10. Few studies have been undertaken on LAB of algal origin and their power  
352 of fermentation of algal extracts. Compared to similar studies, fermentation of lyophilized  
353 commercial *Chlorella vulgaris* microalgae with *Lactiplantibacillus plantarum* and *Lp. pentosus*  
354 resulted in a high decrease of pH until 4.7 after 12h incubation (Carmine 2022). In another  
355 study, *Sargassum* spp. was used for lactic acid production using different LAB strains, resulting  
356 in 1.4 g/L concentration (Sudhakar and Dharani 2022). Acidifying capacity of LAB strains is  
357 essentially based on the conversion of carbohydrates into lactic acid. This conversion is one of  
358 the most important steps for the food industry. To date, lactic acid has been reported to be  
359 produced from starchy and lignocellulosic biomass sources, such as corn and sugar cane waste,  
360 but there are very few reports of lactic acid production from algae (Lin et al. 2020).

361

### 362 **Conclusion**

363 Results of this study showed that marine algae can be considered as an excellent source for the  
364 isolation of new LAB strains potentially endowed with interesting metabolic capacities. The  
365 obtained results showed good performance of LAB isolates to ferment bulk marine algae. The  
366 ability of the isolated strains to ferment algae extracts opens up future perspectives and  
367 biotechnological applications, in particular, the isolation of new antimicrobial substances  
368 (bacteriocins, antifungal compounds, etc.) that could be used as natural biopreservatives in food  
369 industry, extraction of biomaterials (EPS, polylactic acid, etc.) and production of extracellular  
370 enzymes from marine LAB. Finally, isolation of LAB strains from other marine biotopes (other  
371 algae species, sediments, seawater, etc.) and characterization of their properties is needed. The  
372 molecular identification of marine LAB strains also appears very relevant.

373

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379

### 380 **Declaration:**

381

382 **Conflict of interest.** The authors declare that they have no conflict of interest

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