01-May-2017

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Ref: NZJB-2017-0015.R2 Molecular evidence of the presence of Dictyota dichotoma in Argentina.

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Associate Editor, New Zealand Journal of Botany

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- Molecular evidence of the presence of *Dictyota dichotoma* in Argentina based on sequences
   from mtDNA and cpDNA and a discussion of its possible origin
- 3

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## 13 ABSTRACT

14 Dictyota is a brown alga genus inhabiting tropical to warm temperate environments where it 15 performs an important ecological role, such as a food source, shelter and substrate to several 16 species of invertebrates and other algae. The taxonomy of this genus is troublesome, with poor 17 species delimitation and doubtful records in the literature. Dictyota dichotoma, the type species 18 of the genus, was commonly accepted as a cosmopolitan species because of the inaccurate 19 geographical distribution range as a consequence of its misidentification in several parts of the 20 world. Recent studies with molecular data revealed new species and a new understanding of the 21 evolution and biogeography of the genus Dictyota. These studies confined populations of D. 22 dichotoma to Europe (Atlantic and Mediterranean coasts) and the Macaronesian Islands (Azores, 23 Madeira and Canary archipelagos). Also, they confirmed its presence in South Africa, but, 24 whether the species was native or introduced in South Africa could not be verified. In the present 25 study two regions of cpDNA (psbA, rbcL) and one region of mtDNA (nad1) from Argentine 26 samples, identified morphologically as D. dichotoma, were analyzed and compared with other 27 Dictyota species. The identity of these samples as D. dichotoma was confirmed. A haplotype 28 network analysis using all available psbA sequences distinguished seven haplotypes divided into 29 two geographic groups: Atlantic-Mediterranean and Canarian. In Argentina and South Africa, only 30 the most common haplotype of the Atlantic-Mediterranean group was observed. According to the 31 paleoceanographic currents and the presence of European haplotype, the introduced nature of D. 32 dichotoma is indicated.

33

34 Keywords: Dictyotaceae; South Atlantic Ocean; Exotic Species; *psb*A; *rbc*L; *nad*1

35

# 36 INTRODUCTION

*Dictyota* J.V. Lamouroux is an important component of the marine flora in tropical and subtropical environments and is commonly found from the lower range of the intertidal zone to deeper areas (~80m) of the subtidal zone (Littler & Littler 2000; De Clerck et al. 2006; Bittner et al. 2008). The genus has attracted the attention of researchers for several reasons, including the

production of bioactive metabolites (Vallim et al. 2005; De Paula et al. 2011) and its ecological 41 42 role, such as the high biomass that may be used as a food source, shelter and that serves as a 43 substrate to several species of invertebrates and other algae (Genzano & Rodríguez 1998; 44 Stachowicz & Hay 2000; Gauna et al. 2014; Moore & Eastman 2015), as well as the high number 45 of species that make this genus very diverse (Tronholm et al. 2010b). The identification of Dictyota 46 species may be challenging due to phenotypic plasticity, simple morphology and poor delineation 47 between species, which is the reason for the description of hundreds of names for new taxa and 48 doubtful geographical distributional ranges (Schnetter et al. 1987; De Clerck 2003; Tronholm et 49 al. 2010a, 2010b; Gauna et al. 2013). In the southwestern Atlantic, the genus Dictyota remains 50 understudied and the species recorded have never been the subject of a formal review, although 51 there have been some nomenclatural corrections (Oliveira-Filho 1977; Széchy & De-Paula 2016).

52 Dictyota dichotoma (Hudson) J.V. Lamouroux was described as Ulva dichotoma in 53 England in 1762 and later it was transferred to *Dictyota* in 1809 (Lamouroux 1809). During the 54 19th and 20th centuries, D. dichotoma was considered to be widely distributed from tropical to 55 warm temperate areas of the world (Hwang et al. 2005; Tronholm et al. 2008, 2010b) and it 56 was placed in an eurythermic group of species (Van den Hoek 1982). Tronholm et al. (2010b) 57 reviewed the genus Dictyota for Europe using molecular data and a large dataset with sequences 58 from several places of the world and they restricted the occurrence of D. dichotoma to the 59 northeastern Atlantic (Europe and Africa). The specimens from other oceans, previously identified 60 as D. dichotoma, were assigned to other species and so this taxon should not be considered as 61 a cosmopolitan species. So, the native range of Dictyota dichotoma on the European Atlantic 62 coasts was established, from Scotland and southern Norway to Portugal, the Mediterranean Sea 63 and most of the Macaronesia islands, except for the tropical Cape Verde where it was not found 64 (Van den Hoek 1982; Schnetter et al. 1987; Tronholm et al. 2010b). D. dichotoma could not be 65 attributed to natural or introduced origin in South Africa despite that its occurrence there had been questioned by De Clerck (2003), the intriguing record of this species by Tronholm et al. (2010b) 66 67 using a *psbA* marker, and its disjunct distribution.

68 On the Atlantic coast of South America, there are many records of misidentification of 69 species of *Dictyota* as *D. dichotoma* in earlier phycological studies (Saint-Hilaire 1833; Dunal 70 1833; Martius et al. 1833; Montagne 1839). Recently, *D. dichotoma* has been considered as a 71 cryptogenic species in Argentina (Gauna et al. 2014; Raffo et al. 2014; Schwindt et al. 2014).

72 Exotic species are one of the major threats to biodiversity and one of the major concerns 73 throughout the oceans (Schwindt et al. 2014; Abreu et al. 2016). Invasion of marine macroalgae 74 have been reported around the world in recent decades, such as Undaria pinnatifida (Harvey) 75 Suringar, Sargassum muticum (Yendo) Fensholt, Codium fragile ssp. fragile (Suringar) Hariot, 76 Caulerpa taxifolia (M. Vahl) C. Agardh and Schizymenia dubyi (Chauvin ex Duby) J. Agardh 77 (Ramírez et al. 2002; Casas et al. 2004, 2008; Raffo et al. 2009; Irigoyen et al. 2011). Recently, 78 Schwindt et al. (2014) sampled six of the ten main marine ports in Argentina and found the 79 presence of 32 non-indigenous taxa, including exotic and cryptogenic taxa.

This study aims to (i) confirm the molecular identity of specimens previously identified as *Dictyota dichotoma* from Argentina and (ii) discuss the possible origin of the South Atlantic populations.

83

#### 84 MATERIALS AND METHODS

*Dictyota* specimens were collected from Las Grutas, Argentina (San Matías Gulf, 40° 48' S, 64° 48' W) in 2014 from a population previously studied by Gauna et al. (2013, 2014) with respect to the morphology of the specimens, phenology and ecology (epiphytic communities). The algae were collected by SCUBA diving from the subtidal zone and then screened in the field to remove possible epiphytes. Each individual was separated into two sub-samples, one of which was preserved in silica gel for molecular studies, and the other preserved in 4% formalin solution, and deposited at the BBB - Herbarium of Universidad Nacional del Sur (Bahia Blanca, Argentina).

DNA was extracted using HiPuraTM Plant Genomic DNA Miniprep Purification Kit (MolBioTM HIMEDIA). The Plastid-encoded PSII reaction center D1 (*psbA*), NADH *dehydrogenase subunit* 1 (*nad*1) and RUBISCO large Subunit (*rbcL*) were amplified via Polymerase Chain Reaction (PCR) utilizing the primers from Tronholm et al. (2010b) and Bittner et al. (2008) and then sequenced by Macrogen Inc. Korea.

97 The sequences were edited on Mega 6.0 (Tamura et al. 2013) and then analyzed with 98 others retrieved from Genbank (www.ncbi.nlm.nih.gov/genbank) (Table S1). Three datasets were 99 generated with alignments of 695bp for *nad*1, 775bp for *psb*A and 1149bp for *rbc*L. The 100 phylogenetic reconstructions were performed for each gene separately. The concatenation would 101 not be reliable in this case because sequences available on Genbank are from distinct vouchers 102 and from different locations.

103 The evolution model of GTR+I+G was obtained by jModelTest 2.14 for each marker 104 (Darriba et al. 2012) and then used in the Bayesian Inference (BI) analysis on MrBayes 3.1.2 105 (Ronquist et al. 2012). For nad1 and psbA, BI was carried out with 1 million generations in 2 runs 106 and 4 chains, sampling every 1000<sup>th</sup> generation, discarding the first 30 and 50 trees for nad1 and 107 psbA, respectively. For rbcL, BI was carried out with 2 million generations in 2 runs and 4 chains, 108 sampling every 1000th generation, discarding the first 50 trees. The Maximum Likelihood (ML) 109 analysis was performed on Mega 6.0, using the GTR+I+G model with bootstrap of 1000 110 replications.

111 The gene with a higher number of Dictyota dichotoma sequences available from Genbank 112 was psbA and therefore it was used to build a haplotype network with Dnasp v.5 (Librado & Rozas 2009) and Network v.5 by Median Joining. The initial alignment used the 220 available sequences 113 114 (Table S2). Then, the shorter sequences and those with many missing data were removed from 115 the analysis, leaving 149 sequences. To account for the disparity in the number of sequences 116 available from different geographic regions/localities, up to 10 sequences/haplotypes per 117 population were used in the analysis, considering that sequences retrieved from samples 118 collected from up to 100 km apart were considered as part of the same population. The final 119 dataset used in the haplotype network analysis included 68 sequences with 638bp (Table S2).

120

#### 121 RESULTS

122 The topology of the trees for psbA, rbcL and nad1 obtained in the present study agreed 123 with previous studies (Tronholm et al. 2010a, 2010b). The samples from Argentina formed a clade 124 with genuine European sequences of D. dichotoma. For psbA the clade with sequences from 125 Argentina and from the neotype elected by Tronholm et al. (2010b) obtained bootstrap values of 126 100 to ML and 1.00 of posterior probability in the BI analysis (Figure 1). For rbcL (Figure S1) high 127 values of bootstrap and posterior probability were obtained for the clade with sequences from 128 Argentina and Ireland. High values were also observed for nad1 (Figure S2) for the clade with 129 sequences from Argentina and France.

130 The haplotype network (Figure 2) revealed seven haplotypes in the northeast Atlantic 131 ocean that were geographically divided into two groups: Atlantic-Mediterranean and Canarian. 132 The Atlantic-Mediterranean group consisted of two haplotypes, with H1 being the most 133 widespread and present on the Atlantic coasts of the Iberian Peninsula, France, the British Islands 134 and the North Sea, the entire Mediterranean Sea and part of the Macaronesia Islands (Azores 135 and Madeira Archipelagos). The second haplotype (H2) seems to be less frequent in general and 136 was only detected in the Gulf of Lion (France), in the Mediterranean Sea. Samples from Argentina 137 and South Africa corresponded to the H1 haplotype. The Canarian group consisted of five 138 haplotypes (H3, H4, H5, H6 and H7) that are almost restricted to Macaronesia (Canary and Madeira Islands), whereas H5 is the most common one, widely found throughout these islands. 139 140 The H3 haplotype from the Canarian group is the only one that was detected away from these 141 islands in the Gulf of Lion.

142

#### 143 DISCUSSION

144 The higher haplotype diversity observed for *D. dichotoma* in the northeastern Atlantic 145 (Figure 2) is consistent with the hypothesis presented by Tronholm et al. (2010b) explaining the 146 low diversity of the genus Dictyota in Europe. This hypothesis states that following the desiccation 147 of the Mediterranean Sea, which caused the extinction of most of the marine biota during the 148 Messinian salinity crisis (6.8 - 5.3 Ma), the Mediterranean Sea was recolonized by Atlantic species 149 from adjacent areas, such the Macaronesian Islands and the northwestern African coast when it was re-flooded (Coll et al. 2010; Tronholm et al. 2010b). The results from the present study 150 151 demonstrated that European Atlantic and Macaronesian populations of D. dichotoma were 152 probably separated during this vicariant event and, later, northern populations (haplotype H1) 153 were successful in recolonizing the Mediterranean Sea and subsequently reached Macaronesia 154 (Azores and Madeira Archipelagos). Despite having a greater diversity, the Canarian group 155 remained restricted to Macaronesia with the only exception of the haplotype H3, in contrast to 156 what seems to have happened with other Dictyota species from Macaronesia that were able to 157 recolonize the Mediterranean Sea (Tronholm et al. 2010b). However, the presence of the H1 158 haplotype in the South Atlantic would contradict this scenario.

159 The results of Tronholm et al. (2012), using the relaxed molecular clock for the genus, 160 revealed D. dichotoma as the only extant species of one of the oldest lineages in the genus, that 161 split early from the others circa 44.37 Ma (53-34 Ma) and being one of the first clades to disperse 162 through the Tethys Seaway to the westernmost part of the Tethys realm, where it most likely 163 evolved and, later dispersed to the Atlantic coast of Europe and Macaronesia. Since the opening 164 of the South Atlantic (100 - 80 Ma) the northward paleocurrents along the African coast allowed 165 the water transport of the Tethys sea to the Atlantic via Southern Africa while most of the transport 166 continued on account of the circum-equatorial circulation directly to the North Atlantic until 12-18 167 myr (Stille 1992; Stille et al. 1996; Cowman 2013), when the circulation in the North Atlantic was 168 governed by the Pacific-North Atlantic current (Iturralde-Vinent 2006). Therefore, the colonization 169 of the South Atlantic coast of Africa by D. dichotoma would have been prevented because it would 170 have been against the direction of the ocean currents.

171 Furthermore, there was no connection between the westernmost part of Tethys realm 172 and South America, thereby a warm paleocurrent (probably originating in the Caribbean) crossed 173 the Atlantic coast of South America to Patagonia and Tierra del Fuego, where the marine 174 communities were tropical until middle-late Miocene (Del Río 2004a, 2004b; Le Roux 2012). 175 The hypotheses of trans-oceanic dispersal in this scenario fail mainly due to biology of D. 176 dichotoma because culture experiments demonstrated that this species does not possess 177 affinities to tropical conditions (Biebl 1959; Bogaert et al. 2016) which would be required for it to 178 be successfully dispersed along the South American coast until the late Miocene. The adequate 179 temperate marine conditions for D. dichotoma in the southwestern Atlantic were only established 180 after the complete development of the Circumpolar Antarctic Current which lead to the full 181 operation of the Malvinas/Falklands Current and the establishment of the Benguela Upwelling 182 System in the Middle-Late Miocene (10 - 9 Ma). As a consequence, there was a decrease in the 183 seawater temperature in Patagonia and along the southwestern African coast (Heinrich et al. 184 2011; Rommerskirchen et al. 2011; Le Roux 2012), the extinction of the Patagonian tropical 185 marine communities and the retraction of the warm Brazilian current to the north of 186 Argentina/Uruguay (Del Río 2004a, 2004b). Therefore, it is difficult to explain how D. dichotoma 187 would have reached the coast of Argentina through natural dispersion.

188 Those facts agree with the current data that do not show any exclusive haplotypes in 189 the South Atlantic populations (Argentina and South Africa) of D. dichotoma, which would be 190 expected in the case of an old dispersal from the northeastern Atlantic (over 10 myr), or any 191 haplotype shared with the Canarian group, which would be expected in the case of a recent 192 dispersal (less than 6 myr) and so, against the direction of the ocean currents. The occurrence of 193 the H1 haplotype in the South Atlantic ocean is inconsistent with the hypothesis of natural 194 dispersal with subsequent genetic differentiation and supports the hypothesis of human-mediated 195 introduction, where propagules of *D. dichotoma* settled successfully in Argentina and in South 196 Africa because both areas belong to temperate provinces, similar to its native area in the 197 northeastern Atlantic (Spalding et al. 2007).

198 The first evidence of the presence of D. dichotoma in Argentina based on 199 chemosystematics, rather than just morphology, is present by Palermo et al. (1994) who studied 200 a population from Nuevo Gulf and identified three prenylated diterpenes expected to occur (and 201 its precursors) in this species (Amico et al. 1976; Fattorusso et al. 1976; Faulkner et al. 1977; 202 Siamopoulou et al. 2004; Vallim et al. 2005). The locations where D. dichotoma was collected by 203 Palermo et al. (1994) for this study are near to two of the six main marine ports studied 204 by Schwindt et al. (2014). These authors demonstrate that the collection areas are in natural bays 205 with anthropic influences (but with abiotic conditions adequate for Dictyota dichotoma) and have 206 high maritime activities, which enable the high percentage of non-indigenous marine taxa found 207 in them. Port areas provide artificial structures that favor the introduction of exotic fouling/benthic 208 species (recruitment, survival and dispersal) which are mainly transported by ballast water 209 (Schwindt et al. 2014; Abreu et al. 2016; Lin & Zhan 2016). In South Africa most of the introduced 210 species reported in both the cool and warm temperate provinces are from the Northern 211 hemisphere (65%). At the Agulhas ecoregion, where the presence of D. dichotoma was confirmed 212 (Tronholm et al., 2010), 73 non-indigenous (alien, invasive or cryptogenic) taxa have been 213 reported (Mead et al. 2011; Robinson 2015).

Introduction of exotic species may be overlooked for decades (Abreu et al. 2016),
especially when a group (such as *Dictyota*) has not been formally revised. It is not clear at present
how *D. dichotoma* has affected the marine species in Argentina, especially on the northern parts
of the Patagonian coast where the genus *Dictyota* (reported as *D. dichotoma*) is abundant (Casas
et al. 2004; Gauna et al. 2014). For example, in the Gulf of San José and nearby areas (e.g.
Nuevo Gulf and San Matias Gulf) *D. dichotoma* may cover up to 30% of the entire area at depths
up to 10 meters (Boraso de Zaixso & Zaixso 2007; A. Boraso pers. com.).

221 In the same way, the impact on native *Dictyota* species could not be estimated. The 222 known distribution of the genus Dictyota in Argentina (Figure S3) covers the final part of the Warm 223 Temperate Southwestern Atlantic province and the Atlantic part of the Magallanic province 224 (Spalding et al. 2007). The persistent misidentification for almost two centuries (since Montagne 225 1839) resulted in D. dichotoma being considered as the only species in Argentina and the reports 226 of other species, such as Canistrocarpus cervicornis (as Dictyota cervicornis), D. divaricata and 227 D. dichotoma var. intricata (Taylor 1939; Asensi 1966; Van den Hoek 1982; Boraso de Zaixso 228 1995; Mendoza & Nizovoy 2000; Piriz et al. 2003; Boraso & Zaixo 2012; Croce et al. 2015), were 229 considered as representing a wide morphological variation of it (Boraso 2012). Although the 230 genus has also been found on the Beagle Channel (A. Boraso, pers. com.), Mystikou et al. (2016) 231 published the southernmost records of Dictyota in the southwestern Atlantic. Their molecular data 232 confirmed that the species are distinct from *D. dichotoma* (as also verified in Figure 1), which is 233 the first step towards uncovering the diversity of the genus in Argentina.

The spread of *D. dichotoma* northward along the southwestern Atlantic from the Argentine populations is unexpected because of (i) the Confluence Zone of the southward warm Brazilian current and northward cold Malvinas/Falklands current - that spans from about 25°S to 45°S-, where the water masses are reflected eastward as a South Atlantic current (approximate 238 average axis at 39°S, Bisbal 1995), and (ii) the La Plata river (at 35°S), which results in brackish 239 and turbid water, as well as the lack of a suitable substrate that prevents the occurrence of 240 seagrass and marine benthic macroalgae in the La Plata estuary region, which is the reason for 241 the depauperate flora on the Argentine, Uruguayan and Brazilian coasts under its influence (Coll 242 & Oliveira 1999; Calliari et al. 2003; Acha et al. 2008; Braga et al. 2008; Campos et al. 2008). 243 Also, the only Uruguayan record of Dictyota (as D. dichotoma) was guestioned by Coll & Oliveira 244 (1999) because no specimens were found in any of the mentioned herbaria in the original study 245 and no other specimens have been collected subsequently. Therefore, Dictyota species are 246 separated by more than 1200 km between Mar del Plata (Argentina) and Torres (Brazil) due to 247 the lack of any hard substrate (Baptista 1977; Oliveira Filho 1977).

248 In the southeastern Brazil ecoregion, which includes two important Brazilian harbors, there 249 are seasonal coastal upwelling events and the cold (< 18 °C) water masses may reside throughout 250 the year in a deep benthic system (Coelho-Souza et al. 2012). These waters could sustain 251 organisms with cold waters affinities, such as the brown alga Laminaria abyssalis Joly & Oliveira, 252 which occurs below 50 meters (Guimarães et al. 1986). Moreover, the intertidal Jolyna 253 laminarioides Guimarães and Elachistiella leptonematoides Cassano, Yoneshigue-Valentin & 254 Wynne (Valentin 2001; Cassano et al. 2004) only occur during the upwelling period. This can 255 promote a temporary niche for the introduction of species with cold water affinities. There is no 256 evidence, so far, of the presence of D. dichotoma and most of the morphologically similar 257 specimens are D. menstrualis (unpublished data). Recent reports of D. dichotoma and D. 258 dichotoma var. intricata by Villaça et al. (2010) and Crespo et al. (2014) are due to the citation of 259 old studies

260 In conclusion, the natural dispersal of the species from the northeast Atlantic to Argentina 261 and South Africa would be unlikely, as this temperate species would have to cross the whole 262 equatorial and tropical Atlantic ocean against the direction of the currrent. The occurrence of a 263 single, and the most common, haplotype of the Atlantic-Mediterranean group in the South Atlantic 264 populations suggests that D. dichotoma was introduced. The human-assisted introduction of other 265 taxa of Dictyotaceae has already been reported elsewhere, such as the cases of Rugulopteryx 266 okamurae (E.Y. Dawson) I.K. Hwang, W.J. Lee & H.S. Kim (Verlaque et al. 2009), Dictyota 267 cyanoloma Tronholm, De Clerck, A. Gómez-Garreta & Rull Lluc (Tronholm et al. 2010b; García 268 et al. 2016) and Dictyota furcellata (C.Agardh) Greville (Nelson & Wilcox 2010). Future studies 269 including a more comprehensive sampling are necessary and may reveal whether the introduction 270 of D. dichotoma has occurred once or multiple times, its actual geographical range in Argentina 271 and how it has affected the benthic marine community

272

#### 273 SUPPLEMENTARY MATERIAL

Figure S1: Phylogenetic tree based on *rbc*L sequences, presenting a consensus topology estimated by Maximum likelihood (ML) and Bayesian Inference (BI) analyses.

Figure S2: Phylogenetic tree based on *nad*<sup>1</sup> sequences, presenting a consensus topology
estimated by Maximum likelihood (ML) and Bayesian Inference (BI) analyses.

- 278 Figure S3: Localities where *Dictyota* has been reported from Argentina, including the populations
- 279 of *D. dichotoma* discussed in this paper
- 280 Table S1: Sequences used on the phylogenetic analyses
- 281 Table S2: Sequences used on haplotype network of psbA from Dictyota dichotoma
- 282

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(Argentina) with implications for legislation and monitoring programs. Mar Environ Res.

- 499 where the sequences of *D. dichotoma* on GenBank come from. A) Haplotype network of psbA
- 500 showing the two groups: Atlantic-Mediterranean and Canarian. B) Distribution of the seven
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- haplotypes in the North Atlantic population C) South Atlantic populations with the single haplotype
- 502 H1.

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