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## Seasonal Recruitment and Survival Strategies of Palisada Cervicornis Comb. Nov. (Ceramiales, Rhodophyta) in Coral Reefs

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1	SEASONAL RECRUITMENT AND SURVIVAL STRATEGIES OF PALISADA
2	CERVICORNIS COMB. NOV. (CERAMIALES, RHODOPHYTA) IN CORAL REEFS <sup>1</sup>
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23 Núcleo de Pesquisa em Ficologia, Instituto de Botânica, Av. Miguel Estéfano, 3687 – 04301-902 24 São Paulo, Brazil 25 <sup>1</sup> 23 February 2017 26 27 <sup>2</sup>Author for correspondence: email colladol@fiu.edu 28 Running title: Survival strategies of *Palisada cervicornis* comb. nov. 29 30 Abstract 31 As marine tropical ecosystems deteriorate and lose biodiversity, their communities are shifting to 32 dominance of a few species, altering ecosystem's functioning and services. Macroalgae are are 33 becoming dominant on coral reefs, and frequently observed outcompeting corals. Turf algal 34 assemblages are the base of energy flow in these systems and one of the most abundant types of 35 macroalgae on coral reefs, but little is known about their biology and diversity. Through 36 molecular and morphological analyses, we established the proper identity of the turf-forming 37 species Laurencia cervicornis, and by studying seasonal recruitment and the impact of 38 herbivorous fishes on its abundance, we describe its survival strategy. The molecular analyses 39 using a total of 45 rbcL gene sequences including eight current genera within the Laurencia 40 complex and two new sequences of L. cervicornis, strongly support the new combination of 41 Palisada cervicornis comb. nov. In addition, a detailed morphological characterization including 42 the description of reproductive structures, is provided. P. cervicornis was seen recruiting in all 43 seasons but was typically in low abundance. Specimens grown on tiles in fish exclosure cages 44 were devoured in less than 4 hours when offered to fishes. Even though many species of the 45 Laurencia complex have chemicals that deter herbivory, species within the genus Palisada lack

46	feeding deterrents and are highly palatable. We suggest that P. cervicornis is a palatable species
47	that seems to survive in the community by obtaining a size-refuge from herbivory within turf
48	communities
49	
50	
51	Key words: Algal recruitment, Florida, herbivory, Laurencia cervicornis, Palisada cervicornis,
52	phylogeny, <i>rbc</i> L gene, Rhodomelaceae, taxonomy, turf algae.
53	
54	Abbreviations: BI, Bayesian inference; bp, base pairs; ML, maximum likelihood; NJ, neighbor
55	joining; <i>rbc</i> L gene, large subunit of the Ribulose 1,5-bisphosphate carboxylase/oxygenase gene.
56	
57	Introduction
58	Tropical coastal ecosystems are exposed to major anthropogenic stressors that are causing a shift
59	in dominance and/or composition of species, altering the ecosystem's diversity and functioning
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69	taxonomic and morphological diversity has resulted in a relatively low taxonomic resolution of
70	most ecological studies. The vast majority of studies reporting shifts of species dominance are
71	carried out using the word 'macroalgae' as their unit of measurement, while other studies use
72	form-functional groups as their measurement unit (Bruno et al. 2009, Suchley et al. 2016), and
73	still fewer studies have reported results at the genus and species level (Burkepile 2009,
74	Ceccarelli et al. 2011, Duran et al. 2016). However, particular macroalgal species can have
75	profound effects on coral-algae interactions; for example, crustose coralline algae (CCA) are
76	known to be an important group of algae related with coral settlement; however, only a few CCA
77	species actually facilitate coral recruitment (Harrington et al. 2004). Moreover, it is expected that
78	future changes in temperature and CO <sub>2</sub> will affect the composition of macroalgae and their
79	interactions with corals due to species-specific physiological tolerances and increasing algal
80	allelopathic strength (Ober et al. 2016, Del Monaco et al. (2017).).
81	Macroalgal turf, a loosely defined assemblage of aggregated compact small algae
82	(Connell et al. 2014 for a review of the term), likely play important roles in coral reef
83	communities, such as primary producers, structure providers, nitrogen fixers, and facilitate the
84	accumulation of sediments (McCook 1999 for a review). Moreover, algal turfs can have both
85	positive and negative effects on coral recruits (Arnold et al. 2010, Venera-Ponton et al. 2011),
86	and through direct contact, turfs can negatively affect coral growth (Wild et al. 2014). Turfs in
87	the Caribbean Sea are composed of species from such genera as: Laurencia J.V. Lamouroux,
88	Sphacelaria Lyngbye, and Gelidium J. V. Lamouroux, and small filamentous species of all four
89	algal phyla and cyanobacteria (Hay 1981, Carpenter 1986, Littler and Littler 2013). In the
90	Florida Keys coral reefs, species of the Laurencia complex are frequently observed, but their
91	abundance falls into the lower end of the dominant groups of algae, while brown algae such as

Dictyota spp., Lobophora spp. and the green calcareous Halimeda spp. occupy more of the reef
(Yñiguez et al. 2015). However, recruitment and successional studies in the area show that
Laurencia species are more abundant than previously recognized (Duran et al. 2016) but difficult
field identification and consistent removal via high grazing pressure limit their recognition.
Furthermore, because Laurencia spp. are turf-forming, their abundance might be frequently
underestimated. Thus, studies may frequently overlook species with potentially important roles
on coral reefs.

99 Regulation of algal abundance has been related to the abundance of herbivorous fishes. 100 Several experimental studies demonstrate that fish grazing can substantially influence the 101 abundance of different macroalgal species (e.g. Bellwood et al. 2006, Blanco et al. 2011). 102 However, algae can avoid grazing by producing biologically active compounds that deter 103 herbivory (Gressler et al. 2010, 2011). Laurencia spp. are some of the most chemically defended 104 seaweeds and produce a large number of secondary metabolites (Pereira et al. 2003, Manilal 105 2011), 400 of which might have some deterrent effect on fishes (Hay et al. 1988). This might 106 explain why herbivory is relatively low for some *Laurencia* species (Loffler et al. 2014). 107 However, some species within the genus contain similar secondary compounds that do not affect 108 grazing (Hay et al. 1988). Furthermore, some species, such as *Laurencia intricata J.V.* 109 Lamouroux, are nutritionally rich in lipids, protein, amino acids and fatty acids (Gressler et al. 110 2010) making them highly palatable. Although species-level identification of algae is difficult 111 and often requires specialized methods that go beyond morphological characterization, proper 112 identification of species provides important information about many characteristics of organisms 113 that clarify their role in the ecosystem (Knowlton and Jackson 1994, De Clerck et al. 2013, Fong 114 and Fong 2014).

115 The taxonomy of the red algal genus *Laurencia* is extremely complicated due to the large 116 degree of morphological plasticity, the worldwide distribution from temperate to tropical oceans, 117 and the diversity of environments in which it is observed (Fujii et al. 2011). Consequently, the 118 taxonomic position of species within the *Laurencia* complex has rapidly changed as new 119 morphological and molecular data are recognized. Over the past two hundred years, since the 120 establishment of the genus Laurencia (Lamouroux 1813), many taxonomic changes have been 121 proposed resulting in the current Laurencia complex. Currently, it is composed of eight 122 formally proposed genera: Laurencia J.V. Lamouroux sensu stricto, Osmundea Stackhouse 123 (1809), Chondrophycus (Tokida & Y. Saito) Garbary & J.T. Harper (Garbary and Harper 1998), 124 Palisada (Yamada) K. W. Nam (Nam 2007), Yuzurua (K.W. Nam) Martin-Lescanne (Martin-125 Lescanne et al. 2010), Laurenciella Cassano, Gil-Rodríguez, Sentíes, Díaz-Larrea, M. C. 126 Oliveira & M. T. Fujii (Cassano et al. 2012a), Coronaphycus Metti (Metti et al. 2015), and the 127 most recently established Ohelopapa F. Rousseau, Martin-Lescanne, Payri & L. Le Gall 128 (Rousseau et al. 2017). Within this complicated taxonomic reality, it is difficult to come up with 129 proper identification of this important complex of species on coral reefs, causing potential 130 underestimation of their diversity and abundance. 131 In this study, taxonomists and ecologists joined forces to properly identify a turf-forming 132 species in coral reefs, and understand the causes of its low abundance. During a recruitment and 133 successional study of macroalgae in the Florida Keys (Duran et al. 2016), we noticed the 134 presence of a unique tiny turf-forming species that we identified as *Laurencia cervicornis* 135 Harvey. This tiny (< 1 cm tall) species was previously merged by Howe with *Laurencia* 136 corallopsis (Howe 1918), now known as Palisada corallopsis (Montagne) Sentíes, M.T. Fujii &

137 Díaz-Larrea (Sentíes and Díaz-Larrea 2008); however, Littler and Littler (2000), and Wynne et

138 al. (2005) did not recognize this merger, keeping both species: L. cervicornis and P. corallopsis 139 as taxonomically distinct entities. Furthermore, the previously known Laurencia coelenterata D. 140 L. Ballantine & Aponte, now established as Osmundea coelenterata (D. L. Ballantine & Aponte) 141 M. T. Fujii, Sentíes & Areces (Fujii et al. 2016), is also a small species that in the field can easily 142 be confused with L. cervicornis or P. corallopsis unless clear iridescence is observed. We were 143 able to identify and study this species only by growing it on tiles kept in the laboratory under 144 controlled conditions in the total absence of grazers. Therefore, the first goal of this study was to 145 determine the taxonomic identity of our specimens and provide a detailed morphological and 146 molecular analysis of this species. Secondly, to understand the forces controlling the extremely 147 low abundance of Laurencia cervicornis in the Florida Keys, we asked the following questions: 148 1) Is L. cervicornis regularly recruiting? And if so, how frequent is this species recruiting in the 149 study site? 2) If recruitment is frequent, why is the species not abundantly present? We expected 150 to find recruits of *L. cervicornis* throughout the year, but its abundance would be rapidly 151 controlled by herbivores, which could explain the extremely low abundance of this species in the 152 field.

153

154 Materials and Methods

155 *Study site* 

156 This study was conducted in the Florida Reef Track, near Pickles Reef (Key Largo,

157 Florida 25°00'05" N, 80°24'55" W) in a spur and groove reef at a mid-depth area (5-6 m).

158 Parrotfish and surgeonfish are the dominant herbivorous fishes on Pickles Reef with an average

abundance of 5087.17 g per 100 m<sup>2</sup> while the long-spined urchin, *Diadema antillarum* Philippi

160 (1845), is present, only at very low densities (<1 individual per 50 m<sup>2</sup>) (Duran et al. 2016). Water

161 temperature varies seasonally ranging from 24 °C in winter (December and January) to 30 °C

162 during summer. Collection of material and experiments were conducted with the approval of the

163 Florida Keys National Marine Sanctuary (permits: FKNMS-2009-047 and FKNMS-2011-090).

164 *Taxonomic approach* 

Samples of *Laurencia cervicornis* were collected from Pickles Reef, Key Largo, Florida,
USA (Table S1).

167 Molecular analyses

168 The samples used for molecular analysis were dried in silica gel. The total DNA was 169 extracted, using the DNeasy Plant Mini Kit (Qiagen, Valencia, CA, USA) according to the 170 manufacturer's instructions. DNA was amplified by polymerase chain reaction (PCR) using the 171 reaction mix of the Promega® (Madison, WI, USA) in a final volume of 25 µL. The samples 172 were amplified in three overlapping parts with the primer pairs: FrbcLstart - R753, F492 - R1150 173 and F993 - RrbcS (Freshwater and Rueness 1994). Successfully amplified products were purified 174 with the column MicroSpin<sup>™</sup> S-300 HR (GE Healthcare, Buckinghamshire, UK) following the 175 manufacturer's protocol. Sequencing reactions were performed with BigDye<sup>TM</sup> Terminator v3.1 176 Cycle Sequencing Kit (Applied Biosystems, Foster City, USA) on an ABI PRISM 3100 Genetic 177 Analyzer (Applied Biosystems). The primers used for sequencing were the same used for the 178 PCR amplification. 179 Sequence alignments and phylogenetic analysis 180 DNA sequences were analyzed and manually edited in BioEdit v7.0.9.0 (Hall 1999). 181 Identity of generated sequences was checked through BLAST available at the NCBI website

182 (http://www.ncbi.nlm.nih.gov).

183 The model used in the Bayesian inference (BI) and maximum likelihood (ML) analyses 184 was the general-time-reversible model of nucleotide substitution with invariant sites and gamma 185 distributed rates for the variable sites (GTR + I + G). This model was selected using jModeltest 186 2.1.10 (Darriba et al. 2012) under the Akaike information criterion (AIC) as implemented on the 187 CIPRES Science Gateway v3.3. For Bayesian inference analysis, two runs of four chains of the 188 Markov chain Monte Carlo (one hot and three cold) of 10,000,000 generations, with sampling 189 every 1,000 generations, and the initial 10,000 generations in both runs were discarded as 'burn 190 in' to build the consensus tree using MrBayes (v3.2) (Ronquist et al. 2012) implemented on the 191 CIPRES portal. Maximum Likelihood (ML) was performed with PhyML (Guindon and Gascuel 192 2003) using TOPALi v2.5 graphical interface (Milne et al. 2004) under heuristic search with 100 193 bootstrap replicates. Neighbor-Joining (NJ) analysis was performed under heuristic search with 194 2,000 bootstrap replicates using PAUP 4.0b8 (Swofford 2002). Pairwise distances were 195 calculated using the uncorrected 'p' distances in PAUP.

196 Morphology

197 Voucher specimens and material for morphological studies were fixed in 4% 198 formalin/seawater solution or pressed as herbarium sheets. Transverse and longitudinal hand-199 sections were obtained with a razor blade and stained with 0.5% aqueous aniline blue solution 200 acidified with dilute 1N HCl. The microscopic measurements were obtained from the middle 201 portions of the thallus using a calibrated ocular micrometer. Photomicrographs were obtained 202 using a Zeiss Axiocam ERC-5S digital camera (Göttingen, Germany) coupled to an Axio skop 2 203 Zeiss microscope (Göttingen, Germany) and Stemi SV6 Zeiss stereomicroscope (Göttingen, 204 Germany). The vouchers were deposited in the herbaria of the Botanical Institute, São Paulo 205 (SP), Brazil, of the University of São Paulo (SPF), and Fairchild Tropical Botanical Garden

- 206 Herbarium (FTG 167768). Abbreviations follow on-line Index Herbariorum:
- 207 http://sciweb.nybg.org/science2/IndexHerbariorum.asp (Thiers 2017, continuously updated).
  208

209 Ecological approach

210 Recruitment experiment

Eight experimental plots (9 m<sup>2</sup>:  $3 \times 3 \text{ m}$ ) were established in June 2009 to examine the 211 212 effects of herbivory on benthic community dynamics. Briefly, within each plot we established 2 213 1 x 1m herbivore exclusion cages and 2 exclusion cage controls made of three walls to mimic the 214 effects on flow but an open top to allow herbivores access (for a detailed explanation of methods 215 see Zaneveld et al. 2016). Recruitment of *Laurencia cervicornis* was assessed using 10 x 10 cm 216 limestone tiles (obtained from quarried South Florida Pleistocene limestone) attached to plastic 217 mesh with plastic cable ties and secured to the ground with galvanized staples. Two tiles were 218 placed in each exclosure cage (n = 8 plots x 2 exclusion cages x 2 exclusion control cages = 32 219 tiles) and left in the field for three months. To assess seasonal variability, tiles were collected and 220 replaced with new tiles every three months. A total of three deployments covering three seasons: 221 fall (September - December 2011), winter (December 2011 - March 2012) and spring (March -222 June, 2012) were accomplished. Collected tiles were brought to the laboratory and placed in 223 individual aquaria to allow algae to grow to a size that allowed species identification. Aquaria 224 were previously prepared to replicate optimal conditions (12:12 light-dark period, salinity: 35-36 225 PSU, temperature: 25-28 °C, constant water circulation and air pump). Within a week of 226 collection, the percent cover of L. cervicornis was visually quantified and sorted out into the 227 following categories: 0.1 % for a single individual occupying <0.5% of the tile; 0.5% for less 228 than three sparse individuals that occupied <1% cover; 1% for >1 individuals that occupied <5%

cover; and then multiples of 5 were used from 5 to 100% coverage. After inspection, tiles were
returned to their corresponding aquaria and maintained for three more months to promote growth
and to discover reproductive structures in the recruited specimens.

The average percent cover of the two tiles placed within each cage for the eight plots was

used to calculate seasonal abundance and compared across seasons using ANOVA. All analyses

were conducted using R program from R Development Core Team (2012), version 3.2.2.

235 Impact of herbivorous fishes on Laurencia cervicornis experiment

236 Feeding assays were used to assess herbivory impact on the abundance of *L. cervicornis*. 237 In May 2015, 50 limestone tiles (similar to those described above) were deployed for 6 months 238 within a single 2 x 1 x 0.5 m (length x width x height, mesh size 2.5 cm diameter) cage at a 12 m 239 depth to study recruitment and growth of macroalgae, including L. cervicornis. The cage was 240 located off Key Largo (24.9500° N, 80.4540° W), approximately 50 m from the Aquarius Reef 241 Base. From November 9-13 of 2015, nine tiles per day were removed from the exclosure and 242 exposed to herbivorous fishes each day. Divers surveyed and photographed each tile at 4-hour 243 intervals. The first photographs were taken as soon as the tiles were removed from the cage and 244 were employed as a baseline in the morning at  $\sim 08:00$  (initial t = 0). Additional photographs 245 were taken midday ( $\sim$ 12:00; t = +4 hours) and late afternoon ( $\sim$ 16:00; t = +8 hours). We 246 calculated the percent cover of L. cervicornis from photos at each time point using Vidana spatial 247 ecology software (http://www.marinespatialecologylab.org/resources/vidana/). Percent cover was 248 averaged within time points (initial, noon and evening) and repeated measurement ANOVA was 249 used to compare removal rate of *Laurencia cervicornis* within the treatment day. All analyses 250 were conducted using the R program (R Development Core Team 2012, version 3.2.2).

251

252 Results

*Laurencia cervicornis* is a tiny iridescent species living intermingled with other small turf-forming species. In the study area, turfs were frequently covered by sediments making it difficult to distinguish species. We discovered *L. cervicornis* in recruitment tiles kept in aquaria in the laboratory. In the field, it was necessary to flush the sediments covering turfs in order to detect some small iridescent tips intermingled with the rest of turf-forming species. Accordingly, it was impossible to estimate *L. cervicornis* abundance in the field and the results presented here are restricted to observations from our recruitment tiles.

260 Molecular analysis

261 A total of 45 *rbcL* gene sequences were used in this study, including three newly 262 generated sequences, two of L. cervicornis and one of P. corallopsis from Florida, US. The 263 remaining sequences were obtained from GenBank. Two species of Rhodomelaceae were used 264 as outgroups, Chondria collinsiana M. Howe and C. dasyphylla (Woodward) C. Agardh (Table 265 S1). The final rbcL gene alignment consisted of 1,446 bp. The topology of the consensus tree is 266 shown in Fig. 1. The Laurencia complex was resolved as monophyletic with full support based 267 on the models applied to the sequences analyzed. Species of the Laurencia complex were 268 separated into eight clades corresponding to the current genera of the complex: *Chondrophycus*, 269 Coronaphycus, Laurencia, Laurenciella, Palisada, Osmundea, Ohelopapa and Yuzurua, with 270 high to moderate support values, except Ohelopapa that joined the Palisada but without support. 271 Within the *Palisada* assemblage, two main subclades were recognized, both with high support. 272 The two samples of *L. cervicornis* grouped with full support and the intraspecific divergence was 273 0.2%. L. cervicornis is closest phylogenetically to P. furcata and P. corallopsis from which it 274 diverged by 3.1-3.5% and 2.7-3.3%, respectively. Palisada corallopsis from Florida diverged

from the sample from Mexico by 0.4%, indicating that both are the same taxonomic entity. The interspecific divergence within the *Palisada* clade ranged from 1.4% (*P. cf. perforata* and *P. cf. cruciata* from New Caledonia) to 6.8 % (*P. corallopsis* from Mexico and *P. perforata* from the

278 Canary Islands).

279 Our phylogenetic analyses strongly support the transfer of *Laurencia cervicornis* to the genus

280 *Palisada*, and the nomenclatural change is proposed here:

281 *Palisada cervicornis* (Harvey) Collado-Vides, Cassano et M. T. Fujii comb. nov.

282 Basionym: Laurencia cervicornis Harvey, Smithsonian Contributions to Knowledge 5(5): 73, pl.

283 18C, 1853.

284 Morphology

285 Plants grown in aquaria were erect, forming reddish brown tufts or hemispherical clumps 286 up to 5-6 cm high but were never observed over 1 cm high in the field. The plant exhibits blue 287 iridescent rings throughout the thalli (Fig. 2A). The thalli are terete, cartilaginous in texture, and 288 are 0.7-1.2 mm in diameter. Erect axes arise from a single discoid holdfast. Upright branches are 289 scarcely ramified; branching is irregular to dichotomous usually with 1-2 orders of branches 290 (Fig. 2A). Cells in surface view are polygonal, isodiametric, 22-48 µm long and 16-34 µm wide. 291 Secondary pit connections between cortical cells are present (Fig. 2B). In the transverse section, 292 the thalli have one or two layers of pigmented cortical cells 17-23 µm long and 19-32 µm in 293 diameter, and four or five layers of colorless medullary cells, rounded or slightly radially 294 elongated, 51-101 µm long and 30-83 µm wide. Medullary cell walls are uniformly thickened, 295 but lenticular thickenings are absent. Each vegetative axial segment produces two pericentral 296 cells (Fig. 2C). Tetrasporangial branchlets are cylindrical, simple or compound, 1.0–2.7 mm long 297 and 0.5-1.2 mm wide. The arrangement of the tetrasporangia is in a right-angle pattern in

298 relation to fertile branchlets. Mature tetrasporangia are tetrahedrally divided, 38-91 um in 299 diameter. In female thalli, cystocarps are conical with a protuberant ostiole, developed at the 300 subapical portions, partly immersed in the branches, 578-852 µm in diameter. Carposporangia 301 are clavate, 78-113 µm long and 24-95 µm in diameter (Fig. 2D). Male branches are 302 characteristically swollen, 0.5-1.2 mm in diameter. In longitudinal section through a fertile 303 branchlet, the spermatangial pits are cup shaped, and an axial cell row is discernible at the base 304 (Fig. 2E). Spermatangial trichoblasts arise from axial cells, consisting of fertile and sterile 305 branches (Fig. 2F); the fertile branches produce many ovoid spermatangia, 7-10 µm long and 306 3.5-5.8 µm in diameter, and terminate in vesicular sterile cells, 19-30 µm long and 13-21 µm in 307 diameter; each spermatium possesses an apical nucleus (Fig. 2F).

308

309 Ecological approach

310 Recruitment and seasonality

311 Palisada cervicornis recruited to each of the 32 tiles throughout the study period; no significant

312 differences were found between exclosure and control treatments (ANOVA,  $F_{1,69} = 0.011$ , p =

313 0.917). Seasonal differences in recruitment were significant with highest abundance in spring

314 compared with fall and winter (Fig. 3; ANOVA,  $F_{2,69} = 4.499$ , p = 0.014; Tukey posthoc,, spring

315  $\neq$  (fall = winter), p = 0.013)..

316 *Herbivory impact on* Palisada cervicornis *abundance:* 

317 After six months growing inside an exclosure cage, all 50 recruitment tiles were colonized by *P*.

- 318 *cervicornis*, with an average of 5% percent cover. As soon as the tiles were exposed to fish
- 319 grazing, the abundance of *P. cervicornis* rapidly decreased, with the maximum reduction
- 320 occurring in the first 4 hours of exposure (ANOVA,  $F_{2,111}$  = 43.361, p < 0.0001, Tukey posthoc, p

321 < 0.0001, am  $\neq$  (noon = afternoon)); when total consumption of *P. cervicornis* was observed 322 (Fig. 4).

323

324 Discussion

Discovering a tiny, inconspicuous species during an ecological study provided us with the opportunity to address taxonomic and ecological questions in a combined effort. Using molecular tools, we properly identified the species and readjusted its nomenclature status, and through field and laboratory experiments we identified survival strategies of *Palisada cervicornis* in a Florida coral reef.

330 *Taxonomic approach* 

331 The samples from the Florida Reef Track, clearly grouped in the *Palisada* clade distant 332 from the *Laurencia* and *Chondrophycus* clades. Therefore, the sequences from Florida strongly 333 support the new combination here proposed. Intraspecific divergence was minimal (0.2%) and 334 within the range of other *Laurencia* complex species that had divergences of 0.01% to 0.02% for 335 Palisada poiteaui (Díaz-Larrea et al. 2007), 0%-0.4% for Palisada perforata (Cassano et al. 336 2009), and 0%-0.9% for Laurencia dendroidea (Cassano et al. 2012b). Furthermore, the 337 divergence between P. cervicornis and P. corallopsis (2.7-3.3%) establish these two species as 338 separate taxonomic entities, solving the previously proposed merger of those species (Howe 339 1918). Even though the type locality for *Palisada cervicornis* is Key West and the samples for 340 this study come from the upper Keys, this sequence can be representative for the type locality as 341 it belongs to the Florida Reef Track. As for *P. corallopsis*, this is the first reported sequence for 342 the species in Florida, differing by only 0.4% from the sequence reported for the Mexican Caribbean (Díaz-Larrea et al. 2007). The sequenced sample of *P. corallopsis* is from a region 343

near its type locality, which is Cuba (Howe 1918). Therefore, we consider this sequence as
representative of the species.

346 Although Palisada cervicornis (as Laurencia cervicornis) has been reported in several 347 sites around the world, many of those reports do not include a morphological description (Suárez 348 2005, Wynne et al. 2005, Wynne 2011, Tsuda and Walsh 2013); or the descriptions are for 349 juvenile organisms (Littler and Littler 2000, Dawes and Mathieson 2010). Thanks to the 350 cultivation of the specimens of *P. cervicornis* in aquaria, we were able to describe the 351 morphology, including for the first time a detailed description of its reproductive structures and 352 other observations allowing us to differentiate this species from other similar species in the 353 region.

354 Morphologically, Palisada cervicornis is easily confused with Osmundea coelenterata 355 (originally described as *Laurencia coelenterata*), Yuzurua iridescens (M.J. Wynne & D.L. 356 Ballantine) Sentíes & M.J. Wynne and P. corallopsis due to the compact and small thalli and 357 their turf-forming habit. All four species share morphological features such as, two pericentral 358 cells per vegetative axial segment, and arrangement of the tetrasporangia at a right-angle. 359 Furthermore, three of the four species show secondary pit connections between adjacent cortical 360 cells, with the exception being *P. corallopsis*. However, each one possesses its own distinctive 361 characteristics; for example, Osmundea coelenterata has the filament-type of spermatangial 362 branches originating from cortical cells, and tetrasporangia are cut off randomly from the cortical 363 cells, both typical of Osmundea (Fujii et al. 2016). Yuzurua iridescens possesses all generic 364 features established by Nam (1999, as subgenus Yuzurua) (Sentíes et al. 2015), and differs from 365 P. cervicornis by the presence of cortical cell walls near apices markedly projecting with 366 apiculate tips. P. cervicornis and P. corallopsis both form small turfs, but P. corallopsis presents

367 a height of up to 8 cm and no iridescence. O. coelenterata shows a partial iridescence in the 368 apical section of the branchlets (in situ observations), and very small size (to 4 mm), while P. 369 *cervicornis* is larger (2.5 cm in the field, but grows larger in the tanks) and exhibits iridescence 370 along all branches. An examination of a paratype specimen of O. coelenterata (as L. 371 *coelenterata*) from Puerto Rico (#4551) revealed iridescent rings throughout the thalli, the same 372 as *P. cervicornis*; however, the differences in both species are at the genus level. The results of 373 the molecular analyses, and the in-depth morphological observations obtained in the present 374 study corroborate the current taxonomic position of these closely related species of the 375 Laurencia complex.

376 *Ecological aspects of the turf-forming alga* Palisada cervicornis.

377 Coral reef systems have been characterized as grazing controlled ecosystems (Burkepile 378 et al. 2013), where highly productive algal turfs support large grazer communities (Poulin and 379 Klumpp 1992). In turn, these herbivores often affect the abundance and diversity of the algal 380 species present (Hay 1981, Duran et al. 2016) and their productivity (Carpenter 1986, Russ 381 2013). We show that although *Palisada cervicornis* is inconspicuous, it is a common member of 382 the turf community in the Florida Keys that recruits year-round. However, when allowed to grow 383 in herbivore exclusion cages, large *P. cervicornis* are promptly consumed upon exposure to 384 herbivorous fishes (100% consumption after 4 hours of exposure to grazing). In general, the 385 survival of species within the *Laurencia* complex is attributed to the chemical deterrents the 386 algae produce (Hay et al. 1988, Pereira et al. 2003, Malinal 2011). For example, Laurencia 387 *dendroidea* (as *L. obtusa*) produces elatol, which significantly deters herbivory and has been 388 found in Laurencia species worldwide (Pereira et al. 2003). Importantly, not all species in the 389 *Laurencia* complex show similar chemical characteristics, particularly species of the genus

390 *Palisada*, which lack elatol, as well as terpenes and acetogenins that are present in all *Laurencia* 391 species tested so far (Fujii et al. 2011). Thus, P. cervicornis seems to persist in the community by 392 obtaining a size-refuge from herbivory within turf communities rather than through chemical 393 defenses commonly found in *Laurencia* species (e.g. Carpenter 1986; Verges 2011). 394 Integration 395 Understanding species interactions and their consequences for ecosystem dynamics 396 remains a challenge. On coral reefs, herbivore control of macroalgae has been one of the most 397 important and well-studied interactions, with direct consequences on management and the 398 development of strategies to protect and facilitate the recovery of these important ecosystems 399 (Hughes et al. 2010, Mumby 2009). 400 Through the combined efforts of taxonomists and ecologists, we are able to explain the 401 rapid consumption of what was previously believed to be, a heavily-defended *Laurencia* species. 402 Further, our recruitment experiments showed that *P. cervicornis* recruits year round; thus helping 403 explain how such a palatable species is able to persist in a heavily grazed system. In an era in 404 which biodiversity loss is a major environmental crisis (Rockstrom et al. 2009), our report of a 405 new taxonomic combination and first molecular sequence near the type locality of the species, 406 sheds light on the ecology of *P. cervicornis* and furthers our knowledge of the algal biodiversity 407 on the Florida Reef track.

408

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Legend of Tables and Figures
Table S1. Taxa used in this study for phylogenetic analysis.
FIG. 1. Consensus tree derived from Neighbor-Joining analysis of <i>rbcL</i> gene sequences.
Bootstrap supports for NJ (2000 replicates) and ML (100 replicates) (>60%) and Bayesian
posterior probabilities (>0.6) are shown at the nodes; - indicates lack of support; * indicates full
support (99-100% bootstrap value for NJ/ML, and 1.00 for PP). Taxa marked in bold indicate newly
generated sequences.
FIG. 2. Morphological features of Palisada cervicornis comb. nov. (A) Habit of a specimen,
scale bar = 3 mm. (B) Cortical cells in surface view, showing secondary pit-connections
(arrows), scale bar = 20 $\mu$ m. (C) Transverse section of the upper portion of a branch showing an
axial cell (a) and two pericentral cells (p), scale bar = 25 $\mu$ m. (D) Longitudinal section through a
female branchlet showing immersed cystocarp, scale bar = 100 $\mu$ m. (E) Longitudinal section
through a male branchlet showing spermatangial branches in cup-shaped tips, scale bar =

731	100 $\mu$ m. (F) Detail of spermatangial branches on trichoblast with two laterals, sterile (arrow) and
732	spermatangial (arrowhead) branches on its suprabasal cell (sbt). Note spermatangia with an
733	apical nucleus, scale bar = $25 \ \mu m$ .
734	
735	FIG. 3- Seasonal abundance of Palisada cervicornis in recruitment tiles.
736	

737 FIG. 4- Experimental herbivory impact on *Palisada cervicornis*.