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On sea turtle-associated Craspedostauros (Bacillariophyta), with description of three novel species

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## **On sea turtle-associated Craspedostauros (Bacillariophyta), with description of three novel species**





### COVER LETTER

We declare that our paper *"On sea turtle-associated Craspedostauros* (Bacillariophyta), **with description of three novel species***"* has not been published previously, that it is not under consideration for publication elsewhere, that its publication is approved by all authors and by the responsible authorities where the work was carried out, and that, if accepted, it will not be published elsewhere in the same form, in English or in any other language, including electronically without the written consent of the copyright-holder. We believe that the presented manuscript is a valuable addition to the scientific literature as it describes three novel sea turtle-associated diatom species and provides some additional information about the epizoic diatom diversity and ecology.

We confirm that there are no known conflicts of interest associated with this publication and there has been no significant financial support for this work that could have influenced its outcome.

We confirm that the manuscript has been read and approved by all named authors and that there are no other persons who satisfied the criteria for authorship but are not listed. We further confirm that all of us have approved the order of authors listed in the manuscript.

We understand that the Corresponding Author is the sole contact for the Editorial process (including Editorial Manager and direct communications with the office). She is responsible for communicating with the other authors about progress, submissions of revisions and final approval of proofs. We confirm that we have provided a current, correct email address, which is accessible by the Corresponding Author.

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Yours faithfully, Roksana Majewska, on behalf of all the co-authors

Roksana Majewsda





Running title: Sea turtle-associated *Craspedostauros*

## **ABSTRACT**

 Despite recent advances in the research on sea turtle-associated diatoms, some of the key aspects of the diatom-sea turtle relationship, including compositional and functional features of the epizoic diatom community, remain understudied and poorly understood. The current paper focuses on four species belonging to the primarily marine diatom genus *Craspedostauros* that were observed growing attached to numerous sea turtles and sea turtle-associated barnacles from Croatia and South Africa. Three of the examined taxa, *C. danayanus* sp. nov., *C. legouvelloanus* sp. nov., and *C. macewanii* sp. nov. represent novel species and are described based on morphological and, whenever possible, molecular characteristics. The new taxa exhibit characters not yet observed in other members of the genus, such as the presence of more than two rows of cribrate areolae on the girdle bands, shallow perforated septa, and a complete reduction of the stauros. In addition, *C. alatus*, recently described from museum sea turtle specimens, is reported for the first time from loggerheads rescued in Europe. A 3-gene phylogenetic analysis including DNA sequence data for three sea turtle-associated *Craspedostauros* species and other marine and epizoic diatom taxa indicated that *Craspedostauros* is monophyletic and sister to *Achnanthes*. This study, being based on a large number of samples and animal specimens analysed and using different preservation and processing methods, provides some new insights into the genus ecology and biogeography, and sheds more light on the level of intimacy and permanency in the host-epibiont interaction within the epizoic *Craspedostauros* species.

 **Key index words:** *Craspedostauros*, barnacle, *Chelonibia*, epizoic diatom, leatherback, loggerhead, phylogeny, *Platylepas*, sea turtle

**Abbreviations:** BS, bootstrap support; CRW, Comparative RNA Web; LM, light microscopy; ML,

maximum likelihood; SEM, scanning electron microscopy; SSU, small subunit

## **INTRODUCTION**

 The increased interest in epizoic, and more specifically, sea turtle-associated diatoms has in recent years brought about some significant advances in our understanding of the complex relationships between diatoms and their animal hosts. As indicated by sSeveral studies, indicated that diatom communities inhabiting both the skin and the carapace of marine turtles are composed largely of species not observed on other biotic or abiotic substrata (Frankovich et al. 2015, 2016, Majewska et al. 2015a, 2015b, 2017a, 2017b, Robinson et al. 2016, Azari et al. 2020). These observations further suggest a certain level of host-specific evolutionary adaptations used by diatoms. Although intimate relationships between animals and microbes are common and extensively studied, reports of truly epizoic microalgae are generally rare (Ezenwa et al. 2012, Redford et al. 2012, Apprill 2017). Perhaps due to the fact that ubiquitous photosynthetic organisms, such as diatoms, are not immediately perceived as an essential element of any vertebrate microbiome, these new findings are 83 particularly noteworthy. <u>Based on their high frequency of occurrence and high relative abundances</u> recorded from various sea turtle species and geographical regions, as well as lack of records from other types of substrata, several of the newly described sea turtle-associated diatom taxa are currently believed to be strictly epizoic or even sea turtle-specific. While this may be true, many other diatoms present in the sea turtle samples are likely opportunistic species that attached to 88 biofilm in the later stages of its development While several of the newly described sea turtle- associated diatom taxa are currently believed to be strictly epizoic or even sea turtle-specific based on their high frequency of occurrence and high relative abundances recorded from various sea turtle species and geographical regions, as well as lack of records from other types of substrata, many other diatoms present in the sea turtle samples are likely opportunistic species that attached to biofilm in the later stages of its development (Majewska et al. 2015b, 2017b, 2019a,b). Although opportunistic taxa the latter group often dominates specific epizoic habitats in terms of the species

### Page 7 of 113 **Details 12 Details 1**

 number, they opportunistic taxa rarely reach high relative abundance, which may suggest their lack of some key functional adaptations to the epizoic lifestyle.

 As it has already been proposed, studies on sea turtle-associated diatoms may shed more light on the mechanistic processes of divergence and adaptive evolution of diatoms. Furthermore, provided the close relationship between epizoic diatoms and sea turtles holds up under the scrutiny of increased data sampling, new diatom-based tools may be designed to assess the overall well-being of the host in the future (Robinson et al. 2016). Currently, however, the role of diatoms in the sea turtle microbiome functioning remains unknown. Thus, there is no evidence in the existing data for a relationship between the presence or absence of certain diatom groups and the etiopathology of various sea turtle illnesses and disorders. In addition, the interplay between the host and non-host factors influencing the epizoic diatom communities is poorly understood. Therefore, before this endeavour can be accomplished, baseline compositional and ecological data on sea turtle-associated 107 diatom flora must be collected.

 The present study focuses on the sea turtle-associated species belonging to the diatom genus *Craspedostauros* E.J.Cox. At present, the genus comprises ten validly described species including one, *C. alatus* Majewska et Ashworth, described from museum specimens of sea turtles (Cox 1999, Sabbe et al. 2003, Van de Vijver et al. 2012, Ashworth et al. 2017, Majewska et al. 2018). *Craspedostauros* is a predominantly marine genus, although *C. laevissimus* (W. et G.S. West) Sabbe is described as "a widespread endemic species restricted to the Antarctic Continent" and may be of brackish or freshwater origin (Sabbe et al. 2003, Van de Vijver et al. 2012). Most of the *Craspedostauros* members share the typical of the genus morphological characters such as cribrate areolae, numerous doubly-perforated girdle bands, two fore and aft chloroplasts, and a usually narrow stauros. Nevertheless, the latter is reduced or strongly reduced in two species: *C. alyoubii* J.Sabir et Ashworth and *C. paradoxus*\* Ashworth et Lobban. Molecular phylogenetic analysis indicated that the genus is closely related to *Achnanthes* Bory and *Staurotropis* Paddock (Ashworth

 et al. 2017). Both taxa, as well as another marine genus *Druehlago* Lobban et Ashworth, which has yet to be characterized molecularly, share several morphological similarities with *Craspedostauros* (Cox 1999, Ashworth et al. 2017). For example, all the above-mentioned taxa possess valves and girdle bands perforated by cribrate areolae. Moreover, *Craspedostauros* and *Druehlago* share the general frustule morphology, including frustules with central constriction (Ashworth et al. 2017), whereas the fore and aft arrangement of chloroplasts, typical of *Craspedostauros,* can be observed in several *Achnanthes* species (Cox 1999). Three novel species, *C. danayanus* Majewska et Ashworth sp. nov., *C. legouvelloanus* Majewska et Bosak sp. nov., and *C. macewanii* Majewska et Ashworth sp. nov., were found in the course of the

 small population of *C. alatus* is for the first time reported from Europe. A large number of samples analysed and different preservation and processing techniques applied allowed us to document the ultrastructure of the frustule and, whenever possible, the morphology of the plastids as well as the colony type and attachment mode of the cells. These observations were supplemented by a 3-gene phylogenetic analysis including DNA sequence data for three sea turtle-associated *Craspedostauros* species and other marine and epizoic diatom taxa.

ongoing survey on sea turtle-associated diatoms and are described in the current paper. Moreover, a

 \* the specific epithet in *Craspedostauros paradoxa* should be changed to '*paradoxus*' following the recommendations of the International Code of Nomenclature for algae, fungi, and plants (Articles 23.5 & 62; Turland et al. 2018).

### **MATERIALS AND METHODS**

*Material collection and preservation*

### Page 9 of 113 and 12 Journal of Phycology

 Diatom samples were collected from captive and wild sea turtles from Croatia and South Africa. All biofilm samples from carapace and skin were taken using single-use sterile toothbrushes according to the sampling protocols suitable for diatom culturing and standard morphology-based diatom analysis proposed by Pinou et al. (2019). In Croatia, 76 (skin and carapace) samples were collected from 38 loggerhead sea turtles *Caretta caretta* L. rescued and rehabilitated at the Marine Turtle Rescue Centre in Aquarium Pula between 2016 and 2019, on the day of or shortly after their arrival at the facility. In South Africa, 196 (skin and carapace) biofilm samples were collected from 78 loggerheads and 20 leatherbacks *Dermochelys coriacea* Vandelli nesting in Kosi Bay (Indian Ocean) over two nesting seasons, in 2017/2018 and 2018/2019. In addition, 6-mm skin biopsy punches were taken from either front or rear flippers of 30 loggerheads and six leatherbacks and preserved in 4 % formaldehyde solution in seawater immediately after collection. Samples of sea turtle-associated barnacles *Chelonibia testudinaria* L. from 100+ loggerheads and *Platylepas coriacea* Monroe et Limpus from 15 leatherbacks were taken using a plastic paint scraper or a blunt knife during four nesting seasons, in 2015/2016, 2016/2017, 2017/2018, and 2018/2019. Barnacle samples comprised of more than one specimen, were divided into two parts and either frozen (- 20°C) or fixed with 4 % formaldehyde solution in seawater. Single-specimen barnacle samples were frozen (-20°C). Furthermore, skin and carapace samples were collected from seven sea turtles (three loggerheads, three green turtles *Chelonia mydas* L., and one hawksbill *Eretmochelys imbricata* L.) resident at the uShaka Sea World in Durban on 28 June 2019. Material collection was performed by, or under close supervision of, qualified field researchers, and the applied techniques and procedures respected ethical principles of the Declaration of Helsinki

(World Medical Association 2013) as well as all applicable national laws.

*Material processing and microscopy*



182 For scanning electron microscopy (SEM), the oxidized suspension was filtered through  $1-\mu m$  or 1.2-µm Isopore™ (Merck Millipore, Darmstadt, Germany) or 3-µm Nucleopore (Nucleopore, Pleasanton, CA, USA) polycarbonate membrane filters. Formalin-preserved skin and barnacle samples were dehydrated in an alcohol series (30%, 50%, 60%, 70%, 80%, 90%, 95%, 99.9%) followed by critical point-drying in an E3100 Critical Point Dryer (Microscience Division, Watford, UK). Subsequently, the samples were mounted on aluminium stubs with carbon tape and sputter- coated with either gold-palladium using Cressington 108Auto and Cressington 208HR sputter- coaters (Cressington Scientific Instruments Ltd., Watford, UK), palladium using a Precision Etching and Coating System, PECS II (Gatan Inc., CA, USA), or iridium using Emitech K575X (Emitech Ltd., Ashford, Kent, UK) and Cressington 208 Bench Top sputter-coaters. Diatom specimens were analysed with JEOL JSM-7800F, JEOL JSM-7001F (JEOL, Tokyo, Japan), FEI



### *Culturing*

 Living diatoms from the fresh material (unpreserved samples containing sea turtle biofilm and filtered seawater; Pinou et al. 2019) were isolated using a glass pipette with a tip pulled and thinned over a flame into 16x100 mm glass culture tubes (South African strains) or plastic culture flasks (Croatian strains) filled with 34 PSU (South African strains) or 38 PSU (Croatian strains) f/2 growth medium (Guillard 1975). Strains were lit by natural light from a south-facing window (South African strains) or white fluorescent light with a photoperiod of 12h (Croatian strains) and maintained at a temperature of 20*–*24°C. The well-growing cultures were divided into two parts, one of which was used for DNA extraction. The remaining part was cleaned with a mixture of 30%  $H_2O_2$  and 70% HNO<sub>3</sub> and rinsed with distilled water until the near-neutral pH of the fluid phase was reached. Croatian strain (PMFTB0003) was cleaned using saturated KMnO<sub>4</sub> solution and ca. 30% HCl following a slightly modified protocol proposed by Simonsen (1974). Permanent microscopy slides and SEM stubs were prepared as described above.

*DNA preparation and phylogenetic analysis*

#### Journal of Phycology **Page 12 of 113**

 The cultures were harvested as cell pellets using an Eppendorf 5415C centrifuge (Eppendorf North America, Hauppauge, NY, USA) for 10 minutes at 8 000 rpm. The QIAGEN DNeasy Plant Mini Kit (QIAGEN Sciences, Valencia, California, USA) was used for DNA extraction following the manufacturer's protocol, with the addition of an initial cell disruption by 1.0 mm glass beads in a Mini-Beadbeater (Biospec Products, Inc, Bartlesville, OK, USA) for 45 sec. PCR-based DNA amplification and di-deoxy Sanger sequencing of small-subunit nuclear rRNA and the chloroplast- encoded rbcL and psbC markers followed Theriot et al. (2010). Phylogenetic analysis of the DNA sequence data was conducted using a three-gene dataset: nuclear- encoded small subunit (SSU) rRNA, and plastid-encoded *rbc*L and *psb*C. Alignment of the SSU sequences, accounting for secondary structure, was done using the SSUalign program (Nawrocki et al. 2009), with the covariance model based on the 10 diatoms included with the program download, plus 23 additional diatoms from the CRW website (Cannone et al. 2002). Post alignment, SSU sequences were concatenated to the chloroplast sequences into a single matrix (Supplementary Table S1). Eight separate partitions were created for the data (SSU paired and unpaired sites, plus the first, second and third codon positions of each of *rbc*L and *psb*C). This dataset and partitioning scheme were run under maximum likelihood (ML) using RAxML ver. 8.2.7 (Stamatakis 2014) compiled as the pthread-AVX version on an Intel i7 based processor, using the GTR+G model. Twenty-five replicates, each with 500 rapid BS replicates, were run with ML optimizations. Bootstrap support was assessed using the BS replicates from the run with the optimal ML score.

## **RESULTS**

*Morphological observations*

*Craspedostauros danayanus* **Majewska & Ashworth sp. nov. (Figs 2–24)**

### Page 13 of 113 and 12 and 13 and 13 bournal of Phycology

 Cells with two fore and aft H-shaped chloroplasts (Figs 2*–*5). Frustules extremely delicate and very lightly silicified (Figs 6*–*16). In girdle view, frustules rectangular, moderately constricted at the centre (Figs 5, 7 & 11). Valves narrow, linear, very slightly constricted in the valve middle, with bluntly rounded apices (Figs 4, 12*–*16).

## *Light microscopy (Figs 12–16):*

 Valve dimensions (*n* = 30): length 28*–*61 μm, width 2–2.5 μm, length/width ratio: 14–30.5. In cleaned (acid-digested) material, partially dissolved valve margins barely noticeable (Figs 14 & 15, arrows), intact frustules absent. Striae indiscernible (Figs 12*–*16). Raphe-sternum thickened, clearly visible (Figs 12*–*16). Raphe straight (Figs 12*–*16). Thickenings at both central and terminal raphe endings (Figs 12*–*16).

## *Scanning electron microscopy (Figs 17–24):*

 Externally: In cleaned material, valve face appearing flat, with very shallow mantle and straight margin (Figs 17 & 18). Striae uniseriate, 49*–*51 in 10 μm, parallel, becoming radiate towards the apices, alternate or opposite, composed of up to eight areolae (Figs 17 & 18). Areolae largely similar in size, becoming somewhat smaller around the central area, squarish to roundish, externally occluded by covered with cribra (Figs 17–19). Each cribrum perforated by 2–8 pores (Fig. s 17–19). Axial area narrow (Figs 17 & 18). Raphe-sternum not raised (Figs 17–19). Raphe branches straight (Fig. 18). Central area large, symmetrical, amygdaliformfusiform (Figs 18 & 19). Central raphe endings straight, elongated, slightly expanded (Figs 18 & 19). Terminal raphe endings disappearing under somewhat triangular silica flaps extending from the raphe-sternum, giving the impression of 261 unilaterally bent terminal raphe fissures (Figs  $17 \& 18$ ). A large, irregular depression present at the apical flap fold (Figs 17 & 18, arrowheads). Shortened striae composed of cribrate areolae radiating

 around the apices beyond the apical silica flaps (Fig. 17). Asymmetrical pore-freehyaline area present beyond the terminal raphe endings in the immediate vicinity of the apical flap fold (Fig. 17). 265 Internally:<sub>5</sub> Rraphe slit opening laterally onto the more or less uniformly thickened and distinctly raised raphe-sternum (Fig. 20). Stauros absent (Figs 20 & 21). Central area mirroring the external 267 structure in size and shape (Figs 20  $\&$  21). Central raphe endings elongated, very slightly 268 unilaterally bent, terminating onto weakly constricted rectelevatum (Figs 20 & 21). Terminal raphe endings positioned somewhat laterally on a large and rounded apical part of the raphe-sternum, terminating in prominent helictoglossae (Figs 20  $\&$  23). Asymmetrical thickening extending from the apical part of the raphe-sternum towards the valve margin, corresponding to the external apical  $2/2$  silica flaps (Fig. 23, arrowheads). Areolae externally occluded with cribra, appearing sunken (Figs 21–23).

 Cingulum composed of numerous (14+) open copulae, bearing two rows of typically squarish, roundish or elongated areolae, ca. 50*–*60 in 10 μm (Figs 18, 23 & 24). Areolae occludedcovered 276 externally by cribra (Figs  $23 \& 24$ ).

### **Taxonomic remarks**

 *Craspedostauros danayanus* is most similar to *C. paradoxus*, sharing the general valve outline and lacking the stauros. However, *C. danayanus* differs from the latter in being distinctly smaller (28*–* 61 μm vs 80*–*85 μm) μm and more slender (2*–*2.5 μm vs 6.5*–*9 μm), possessing a higher stria density (36*–*40 vs 49*–*51 vs 36*–*40), and lacking the lip-like silica flaps (externally) and the central knob (internally) present in *C. paradoxus* (Table 1).

### Page 15 of 113 and 12 and 13 Journal of Phycology



ZA0019A/ZA1824E) deposited in the South African Diatom Collection housed by North-West

University, Potchefstroom, South Africa.

288 TYPE LOCALITY: Mabibi Beach, Elephant Coast, South Africa (27° 21′ 30" S, 32° 44′ 20" E).

Collected from the barnacle *Platylepas coriacea* growing on the egg-lying leatherback sea turtle

(tag numbers: ZA0019A, ZA1824E) by R. Majewska, 7 December 2018.

ETYMOLOGY: The epithet honours Danay A. Stoppel (North-West University, Potchefstroom,

South Africa), who made the first observations of the new taxon, in recognition of her contribution

to the sea turtle diatom project in South Africa.

ECOLOGY: Epizoic on carapaces of adult leatherback sea turtles and on leatherback-associated

barnacles *Platylepas coriacea* growing on adult leatherbacks from Kosi Bay (South Africa).

Attaching to the animal surface through one end of the valve, motile in culture.

 The taxon was found in twelve leatherback skin samples (out of 20 examined) and in all *P. coriacea* 298 samples examined ( $n = 15$ ) reaching relative abundances of 35% (skin samples) and 79% (barnacle samples). It was found in neither loggerhead nor loggerhead-associated barnacle samples from the same location (Kosi Bay, South Africa). Leatherback skin samples containing *C. danayanus* were dominated by *Navicula* spp., *Tursiocola* sp., and *Poulinea* spp. The new taxon was dominant in most of the *P. coriacea* samples along with *Cylindrotheca* sp. Both taxa colonised various anatomical parts of the barnacle showing preference for rough surfaces and cavities. The extremely lightly silicified frustules may be an adaptation to the pelagic lifestyle of the host, as the open ocean waters contain significantly lower concentrations of dissolved silica than coastal habitats (Tréguer et al. 1995).

*Craspedostauros legouvelloanus* **Majewska & Bosak sp. nov. (Figs 25–47)**

### *Light microscopy (Figs 25–30):*

 Intact frustules lying almost always in girdle view (due to large cell depth/valve width ratio), slightly constricted in the middle (Figs 25, 26, 28*–*30), broad with several girdle bands (Figs 26, 28 & 30). Valve margin expanded at the centre (Figs 25, 28 & 30). Frustules lightly silicified and delicate. Valves narrow, linear to linear-lanceolate, slightly constricted at the central area, with bluntly rounded apices (Fig. 27). Valve dimensions (*n* = 30): length 18*–*34 μm, width 3–5 μm, length/width ratio: 5.6–9.4. Striae indiscernible (Figs 25*–*30). Stauros narrow (Figs 25, 27*–*30), widening towards the biarcuate valve margins (Fig. 30, arrows). Raphe-sternum clearly visible (Figs 25*–*30). Raphe straight, biarcuate in girdle view (Figs 25, 26, 28 & 30).

## *Scanning electron microscopy (Figs 318–4017):*

 Externally: Valves somewhat convex, with no clear valve face-mantle junction (Figs 31–336). Valve margin clearly expanded at the centre beyond the stauros (Fig. 33). Striae uniseriate, 46*–*49 in 10 μm, parallel throughout the valve centre, becoming convergent near the apices, alternate or opposite, composed of up to 13 areolae (Figs 31, 32 & 38). Areolae similar in size throughout the entire valve, squarish, externally occluded by evered with cribra (Figs 31–33 & 38). Each cribrum 325 perforated by 4 pores (Figs 31–33 & 38). Axial area very narrow (Figs 31 & –323). Raphe-sternum very slightly raised (Figs 31–33). Raphe branches more or less straight (Fig. 31). Central area forming a narrow rectangular fascia (Figs 31 & 38). Central raphe endings covered entirely by rimmed lip-like silica flaps extending from one side of the axial area (Figs 31 & 38). At the apices, axial area expanding into somewhat triangular silica flaps covering the terminal raphe endings giving the impression of unilaterally bent terminal raphe fissures (Figs 31–33). An oval or irregular depression present at the apical flap fold (Fig. 31, arrows). Shortened stria composed of regular areolae and simple puncta radiating around the apices beyond the terminal raphe endings (Figs 31– 33).

### Page 17 of 113 and 12 and 12 and 13 and 13 and 13 and 13 and 13 and 14 and 15 and 16 Phycology

334 Internally: Rraphe slit opening laterally onto the uniformly thick and clearly raised raphe-sternum (Figs 35 & 36). Stauros raised, very narrow, broadening abruptly at the mantle expansion and merging with the pore-freen value area at the valve margin (Figs 36 & 39), slightly more expanded and somewhat thicker on the side corresponding to the external lip-like silica flaps (Figs 36, arrowheads,  $39 \& 40$ ). Central raphe endings straight or slightly unilaterally bent, elongated, 3<sup>3</sup>89 terminating onto a weakly developed, elongated and rectelevatum flattened helictoglossae (Figs 35, 36, 39 & 40)., Abearing a blunt cylindrical knob with a small central cavity present between the raphe endings (Figs 35, 36, 39 & 40). Areolae externally occluded by espected with cribra, appearing sunken, especially close to the stauros (Figs 39 & 40). Stauros-adjacent virgae appearing hollow, suggesting a more complex valve structure in that area (Fig. 39, arrowheads). Terminal raphe endings positioned somewhat laterally on the raphe-sternum, terminating onto prominent helictoglossae. At the apices, within an expanded raphe-sternum expanded laterally towards the valve margin, merged with pore-freehyaline area corresponding to the external apical silica flaps (Figs 36 & 37).

 Cingulum composed of numerous (12+) open copulae, bearing two rows of typically squarish or elongated areolae, ca. 50*–*60 in 10 μm (Figs 32–35). Areolae occludedcovered externally by cribra with 4*–*12 pores per cribrum (Figs 32–35). Valvocopula curved, distinctly narrower and pore- freehyaline besideat the stauros (Fig. 33, arrowheads). An internal ridgethickening perforated by puncta, resembling a reduced septum, present in each copula except for valvocopula (Figs 33–35, arrowheads).

## *Adriatic population (Figs 41–47)*

 Specimens resembling *C. leguovelloanus* were found on the carapace of six loggerhead sea turtles sampled on the Croatian coast of the Adriatic Sea. Most of the morphological features observed in the Adriatic population (Figs 41–47) agreed well with those found in *C. legouvelloanus* (Figs 41–

### Journal of Phycology **Page 18 of 113**

 47). The cells possessed two fore and aft H-shaped chloroplasts (Fig. 41, arrows) observed previously in other *Craspedostauros* species (Cox 1999, Ashworth et al. 2017, Majewska et al. 2018). The specimens were slightly longer (23*–*39 μm) and wider (3.5*–*6 μm, length/width ratio: 5.2*–*7.8, *n* = 25) than those from the South African population and their stria density was lower (40*–* 44 in 10 μm vs. 46*–*49 in 10 μm; Table 1). In general, the frustules showed a relatively high degree of irregularity in the areolae structure and the size and shape of stauros, axial area, and facia (Figs 42*–*45).

## **Taxonomic remarks**

 Currently, *C. legouvelloanus* is the only *Craspedostauros* species with septate girdle bands. ValvesFrustules of this speciestaxon differ from those of all known stauros-bearing *Craspedostauros* species in possessing a very high stria density (above 40 in 10 μm). Although a similarly high or higher stria density was observed in *C. alyoubii* (~40 in 10 μm) and *C. danayanus* (49*–*51 in 10 μm), the two species are larger (83*–*105 μm and 28*–*61 μm) than *C. legouvelloanus*  (18*–*34 [39] μm) and their general morphology differs remarkably from that of the new taxon in, for example, possessing a reduced or strongly reduced stauros (Table 1). Several of the characters of *C. legouvelloanus*, such as largely uniform valve areolae with four pores per cribrum and internal central knob, agree with the description of *C. australis* E.J.Cox (Cox 1999)*.* However, the new species can be easily distinguished from the latter by its clearly centrally expanded valve margin and well-developed lip-like silica flaps externally covering the central raphe endings absent in *C. australis* (Table 1).

 Although wild specimens belonging to the Adriatic population of *C. legouvelloanus* exhibited numerous irregularities in the shape and size of taxonomically important characters such as areolae, striae, stauros, and central area, we were unable to indicate and unambiguously describe features that would distinguish them from the type population. High morphological plasticity and

### Page 19 of 113 and 113 and 12 Journal of Phycology

 polymorphy in diatoms have been reported from both epizoic and non-epizoic habitats (Cox 2011, De Martino et al. 2011, Urbánková et al. 2016, Riaux-Gobin et al. 2014, 2017, Edlund and Burge 2019), and it is conceivable that the morphological differences observed between the two populations could be induced by environmental triggers, such as differences in salinity or nutrient concentrations (Schultz 1971, Czarnecki 1987, 1994, De Martino et al. 2011). Unfortunately, the Croatian strain PMFTB0003 (Figs 41, 43, 45 & 46) isolated from the sample TB13 did not survive and the DNA material could not be obtained at the time of this study. Therefore, in the light of the current lack of any additional information about the phylogenetic relationships between the two populations, they should be considered conspecific until otherwise proven.

 HOLOTYPE: Permanent slide SANDC-ST003 and unmounted material (prepared from sample ZA0762D/ZA0763D) deposited in the South African Diatom Collection housed by North-West University, Potchefstroom, South Africa.

PARATYPE: Permanent slide HRNDC000150 and unmounted material (TB13) deposited in the

Croatian National Diatom Collection housed by Faculty of Science, University of Zagreb, Croatia.

 ISOTYPES: Permanent slides BR-XXXX and BR-XXXX deposited in the BR-collection housed by Meise Botanic Garden, Meise, Belgium.

401 TYPE LOCALITY: Kosi Bay, South Africa (26° 59' 39" S, 32° 51' 60" E). Collected from the

carapace of the egg-lying loggerhead sea turtle (tag numbers: ZA0762D, ZA0763D) by R.

Majewska, 15 December 2017 (holotype).

404 Marine Turtle Rescue Centre, Pula, Croatia (44°50′ 07" N, 13°49′ 58" E). Collected from a semi- adult female loggerhead *Caretta caretta* named 'Mimi' by K. Gobić Medica, 28 May 2019 (paratype).



## *Craspedostauros macewanii* **Majewska & Ashworth sp. nov. (Figs 48–62)**

## *Light microscopy (Figs 48–54):*

 Cells with two fore and aft H-shaped chloroplasts (Figs 48 & 51). Frustules delicate and lightly silicified (Figs 48*–*54). In girdle view, frustules rectangular, moderately to strongly constricted at the centre (Figs 48*–*50). Cingulum composed of several girdle bands (Figs 498*–*50). Valves narrow, linear to linear-lanceolate, slightly constricted at the central area, with bluntly rounded apices (Figs 51*–*54). Valve margin straight (Fig. 49, arrow). Valve dimensions (*n* = 20): length 26–51 μm (up to  $428 - 65 \mu m$  in culture)  $\mu$ m, width 4.5–5.5  $\mu$ m (up to –6  $\mu$ m in culture)  $\mu$ m, length/width ratio: 5.4–11.3. Valve face-mantle junction visible on each side of the raphe (Figs 52–54, arrows). Striae barely discernible, 28–31 in 10 μm (Figs 52*–*54). Central area narrow, bow tie-shaped (Figs 52*–*54).

### Page 21 of 113 and 12 Journal of Phycology

 Raphe-sternum thickened (Figs 52*–*54). Raphe straight (Fig. 54) with thickenings at the terminal raphe endings (Figs 52*–*54).

## *Scanning electron microscopy (Figs 55–62):*

 Externally: Valves slightly concave at the centre, with distinct valve face-mantle junction marked 486 by a narrow pore-freehyaline area (Figs 55  $&$  57). Valve face flat (Fig. 55). Mantle very deep (Fig.  $4\overline{2}7$  55). Valve margin straight, with narrow pore-free<del>hyaline</del> area at the mantle edge (Figs. 56 & 575). Striae uniseriate, parallel through most of the valve, becoming convergent near the apices, alternate or opposite, composed of up to 21 areolae (2–8 on the valve face and up to 13 on the mantle; Figs 55–58). Areolae similar in size, squarish, externally occludedcovered bywith cribra (Figs 56–58). Areolae bordering the narrow axial area usually only slightly larger and somewhat irregular in shape (Figs 56–58). Each cribrum perforated by highly variable number of pores (up to 13+; Figs 56–58). Raphe branches more or less straight (Fig. 55). Central area in the form of a narrow bow 444 tie-shaped fascia (Figs 55  $\&$  57). Central raphe endings covered by small lip-like silica flaps 445 extending from one side of the axial area (Figs 55 & 57). Apices pore-free hyaline (Figs 55, 56 & 58). Terminal raphe endings covered by triangular silica flaps giving the impression of unilaterally 447 bent terminal raphe fissures (Figs 55, 56 & 58). An oval or irregular depression (Figs 55, arrowhead, 56 & 58) with several small areolae (Figs 56 & 58, arrowheads) present at the apical flap fold. Shortened striae composed of a single areola (occasionally with additional puncta) 450 radiating around the apices beyond the terminal raphe endings (Figs  $56 \& 58$ ). Internally:, Rraphe slit opening more or less centrally onto the uniformly thick raphe-sternum (59– 61). Stauros raised, narrow, tapering towards the valve face-mantle junction and widening 453 significantly on the valve mantle towards the mantle edge (Figs 59  $\&$  61). Central raphe endings

- straight, elongated, terminating onto weakly developed, elongated and flattened helictoglossae
- flattened rectelevatum (Figs 59 & 61). A flatly ended cylindrical knob present at the central nodule

 (Figs 59 & 61). Areolae externally occluded by espected with cribra, appearing sunken, especially 457 close to the raphe-sternum (Figs  $60 \& 61$ ). Terminal raphe endings terminating onto prominent helictoglossae within an expanded and thickened pore-freehyaline area corresponding to the curvature of the external silica flaps (Fig. 60). Several small areolae present at the end of the curved thickening (Fig. 60, arrowheads).

 Cingulum composed of numerous open copulae bearing up to five rows of cribrate squarish or elongated areolae, ca. 38*–*45 in 10 μm (Figs 55, 59 & 62). Advalvar part of valvocopula pore-freehyaline besideat the stauros (Fig. 59).

### **Taxonomic remarks**

 The morphological character pattern in *Craspedostauros macewanii* is most similar to *C. australis*  and *C*. *capensis* Cox. The three species share several features such as the presence of a bow tie- shaped fascia, rudimentary lip-like silica flaps extending from the raphe-sternum and partially covering the external central raphe endings, valve margin straight at the centre, and internally, a single knob at the central nodule (Table 1). Moreover, valve dimensions of *C. macewanii* (26–51 μm long, 4.5–5.5 μm wide) overlap with those reported for *C. australis* (35–78 μm long, 4–6 μm wide) and *C. capensis* (25–35 μm long, 4.5–5.5 μm wide). In *C. macewanii*, however, the stria density (28–31 in 10 μm) is significantly higher than in *C. capensis* (~19 in 10 μm) and lower than in *C. australis* (35 in 10 μm). In addition, *C. macewanii* can be distinguished from both *C. australis* and *C. capensis* by the presence of a distinct valve face-mantle junction running as a narrow, though clearly visible, pore-freehyaline ridge from apex to apex. *Craspedostauros macewanii* differs further from *C. capensis* in possessing areolae of a similar size throughout the entire valve (variable in *C. capensis*), and from *C. australis* in having convergent stria at the apices (parallel in *C. australis*) and extended apical hyaline zone (Cox 1999). The new taxon is also the only



483 HOLOTYPE: Permanent slide SANDC-ST242-and unmounted material (prepared from sample

ST242) deposited in the South African Diatom Collection housed by North-West University,

Potchefstroom, South Africa.

486 TYPE LOCALITY: uShaka Sea World, Durban, South Africa (29° 52′ 02.79″ S, 31° 02′ 45.29″ E).

 Collected from the carapace of a captive juvenile loggerhead named "Bubbles" by R. Majewska, 28 June 2019.

 ETYMOLOGY: The epithet honours Tony McEwan, the uShaka Sea World director, whose scientific enthusiasm and support to the sea turtle diatom project are highly appreciated and acknowledged.

 ECOLOGY: Epizoic on skin and carapaces of captive loggerheads and green turtles. Attaching to the animal surface through one end of the valve, motile in culture.

 The taxon was found on two captive loggerheads (a juvenile named "Bubbles" and an adult female named "DJ") and two captive green turtles (a subadult named "Calypso" and an adult male named

"Napoleon") each time reaching relative abundance of 0.5–1%. All carapace samples containing *C.* 

*macewanii* were dominated by the so-called "marine gomphonemoids": *Poulinea* spp. and

*Chelonicola* spp., accompanied by *Amphora* spp., *Nitzschia* spp., *Achnanthes elongata* and *A.* 

*squaliformis* Majewska et Van de Vijver, whereas the most abundant taxa in the four skin samples

were *Tursiocola* spp., *Medlinella* sp., and the two previously mentioned *Achnanthes* species.

### *Craspedostauros alatus* **Majewska & Ashworth (Figs 63–74)**

 *Craspedostauros alatus* was found on the carapaces of several loggerhead sea turtles sampled at the Marine Turtle Rescue Centre in Pula, Croatia. The taxon co-occurred with *C. legouvelloanus*. As in the case of the latter, relative abundance of *C. alatus* was low (ca. 1–3% of the total diatom number). The observed morphological features of the Adriatic population agreed with the original description of the species (Majewska et al. 2018; Figs 63–74, Table 1). The examined specimens were 26*–*34 μm long and 3*–*5 μm wide (length/width ratio: 6.3*–*8.8), with stria density 24*–*27 in 10 μm (*n* = 20), and possessed all species-specific features, including a very distinct valve face-mantle junction and deep mantle (Figs 68, arrows, 69–71), wing-like silica flaps at the apices (Fig. 70), and 511 rectelevatum with central cavity (Figs 73 & 74).

*DNA-based phylogeny*

 The genus *Craspedostauros* is monophyletic based on DNA sequence data generated from cultured material thus far (Fig. 75), though not with strong bootstrap support (bs < 50%). Regarding the taxa described here, *Craspedostauros macewanii* is sister to the rest of the clade (except *C.* 

 *amphoroides*) with high support (bs = 96%), while *C. danayanus* is sister to *C. alyoubii* and *C. paradoxus* (bs = 71%).

 Consistent with other molecular phylogenetic studies which include the genus (Ashworth et al. 2017), the position of the *Craspedostauros* clade can be found in a poorly supported (bs < 50%) assemblage containing the *Staurotropis* clade and a clade of marine *Achnanthes* species. This assemblage can be found within a clade with the Bacillariales (Supplementary Figure S1), though the relationship between the *Staurotropis*+*Achnanthes*+*Craspedostauros* clade and the three Bacillariales clades is poorly resolved. For taxa, strain voucher ID and GenBank accession numbers for strains used in the analysis see Supplementary Table S1.

### **DISCUSSION**

 The three new species described in the current study share most of the morphological characters typical of the genus *Craspedostauros*, such as squarish or rectangular areolae occluded by cribra on the valve and girdle bands, multiple copulae with at least two rows of perforations, and two fore and aft chloroplasts. Their linear or linear-lanceolate valve outline and the central constriction of the cell seen in girdle view resemble previously described species. Interestingly, two of the novel species, *C. macewanii* and *C. legouvelloanus*, present features not yet observed in any other member of the genus. The former possesses more than two rows of cribrate areolae on the girdle bands, whereas the latter shows shallow perforated septa. Moreover, the leatherback-associated *C. danayanus* presents a complete reduction of the stauros being the second, after *C. paradoxus*, *Craspedostauros* species lacking this character.

 It is interesting to note that as the number of character states, such as the reduction/loss of the stauros (*C. paradoxus* and *C. danayanus*) or addition of septate copulae (C. *legouvelloanus*), within *Craspedostauros* changes, the molecular data remain constant in their support (however tenuous) of monophyly for the genus. Cox (1999) ascribed the constricted girdle view to the presence of stauros. Yet the frustules of the two species lacking the latter, still show the central constriction, which may indicate that the lack of stauros is a secondary loss. One of the morphological features of the genus which has been maintained, regardless of newly described diversity, has been the cribrate areolar covering. While the degree of cribrum poration might change among species, the overall gestalt ultrastructure remains unchanged. Even more interesting is that this cribrum ultrastructure is also seen in *Staurotropis* and the *Achnanthes* species, which are commonly found (again, somewhat tenuously) sister to the *Craspedostauros* clade in molecular phylogenies. While there are other morphological similarities between the three genera, such as the stauros (though missing in some species of *Craspedostauros* and *Achnanthes*) and the fore and aft H-shaped or plate-like chloroplasts (missing in *Staurotropis* and some species of *Achnanthes*), so far it is the cribrate

 areolae ultrastructure that remains constant. In this context, the phylogenetic position of the genus *Druehlago*, which shares the same cribrum ultrastructure and the same chloroplast morphology of *Staurotropis* and *Achnanthes longipes* Agardh, but thus far lacks a stauros-bearing taxon, is all the more intriguing.

 Microscopical analyses of the fresh and critical-point-dried sea turtle skin pieces and barnacles revealed the mode of attachment and growth form of *C. danayanus* that attaches to the animal substratum through one pole of the cell. A similar mode of attachment to the natural substratum was observed in several members of the genus (R.Majewska, pers. observ.) suggesting that these taxa can either develop as firmly attached, sessile colonies or remain motile in less favourable conditions (e.g. in culture tubes).

 In the course of the on-going surveys on sea turtle-associated diatoms, a recently described taxon, *C. alatus*, was observed growing on the carapaces of several loggerhead sea turtles rescued in Croatia. *Craspedostauros alatus* was originally described from museum specimens of juvenile Kemp's ridleys (*Lepidochelys kempii* Garman) and a juvenile green turtle found cold-stunned and beyond recovery on the New York (USA) beaches during various seasons between 2012 and 2014 (Majewska et al. 2018). Although the relative abundance of *C. alatus* did not exceed 5.5% (current study, Majewska et al. 2018), observations of this taxon on a sea turtle from the Adriatic Sea may indicate that a) *C. alatus* is not an uncommon element of the sea turtle diatom flora; b) being associated with highly migratory animals such as sea turtles its geographical range is likely linked to that of its hosts.

 A similar conclusion can be drawn based on the records of *C. legouvelloanus*. The species occurred on several of the Adriatic loggerheads as well as on dozens of sea turtles belonging to the same species and their associated barnacles sampled on the eastern coast of South Africa. Even though the taxon was found in two different ocean basins, it cannot be excluded that the sea turtles acted as vectors that facilitated its dispersal among the various seas and oceans. There is a strong

### Page 27 of 113 and 12 and 13 Journal of Phycology

 observational and molecular evidence that the Indian Ocean loggerheads interact and mate with the Atlantic members of the species (Bowen et al. 1994, Bowen and Karl 2007, Le Gouvello du Timat et al., in prep.). Thus, it is conceivable that any diatom able to endure the changing conditions during the migrations of their hosts and survive in competition with native flora would inoculate all appropriate and available media and substrata encountered. With the exception of *C. danayanus*, the sea turtle-associated *Craspedostauros* species, although common on the sea turtle carapaces, were never among the dominant taxa, and it is still unclear whether the animal body surface is their preferred or alternative habitat. It is possible that the occurrence of these species in the sea turtle biofilm samples is linked to the presence of some other sea turtle epibionts (e.g. barnacles, sponges, bryozoans). *Craspedostauros danayanus* dominated most of the leatherback skin and barnacle samples that were analysed, and it is likely that this taxon is highly adapted to the conditions provided by the smooth body of the largest among the sea turtles, and, being associated with both the skin and the leatherback-specific barnacle species, *Platylepas coriacea*, its relationship with the host may be obligatory. Leatherbacks, contrary to other extant sea turtles, show the fully oceanic developmental pattern spending most of their lives in highly homogenous open-water environment devoid of refugia (Bolten 2003). They are unique among modern reptiles in being endothermal (Frair et al. 1972). This ability allows them to survive in both tropical and near-freezing waters (James et al. 2006). They are also significantly faster swimmers and deeper divers than other sea turtles (Eckert 2002, Doyle et al. 2008). Therefore, microhabitats provided by these animals, and thus their microbiomes, would differ substantially from those present on other sea turtles. Under such unique conditions, far from the diverse, species-rich shallow-water ecosystems, specific eco- physiological adaptations may be required to survive, and fewer diatom species would manage to thrive on the demanding substratum. An analogous phenomenon is known from marine cetaceans that seem to be colonised by only a few, highly specialized diatom taxa (e.g. Nemoto 1956, Holmes et al. 1993, Ferrario et al. 2018).

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## **Figures legends**

 **Fig. 1**. Sampling locations where *Craspedostauros danayanus* (1), *C. legouvelloanus* (2), *C. macewanii* (3), and *C. alatus* (4) were found.

 **Figures 2–11.** *Craspedostauros danayanus*. **Fig. 2.** Living cells of *C. danayanus* and *Cylindrotheca* sp. attached to the leatherback skin scutes (light microscopy). **Fig. 3.** Stained colony of *C. danayanus* and associated bacteria on the leatherback skin scutes. **Fig. 4.** Valve view of a living cell (cultured strain). **Fig. 5.** Girdle view of a living cell (cultured strain). **Figs 6–11.** Scanning electron micrographs of *C. danayanus* attached to its original substratum. **Fig. 6.** Monospecific colony growing among the flaking skin of leatherback (dorsal side of the hind flipper). **Fig. 7.** Extremely delicate and fragile cells of *C. danayanus* attached to the leatherback skin (dorsal side of the hind flipper). **Fig. 8.** An overview of the leatherback-associated barnacle, *Platylepas coriacea*, colonized by *C. danayanus*. **Fig. 9.** A detail of the external part of the barnacle with a sheath of host sea turtle tissue overgrown with *C. danayanus*. Arrows indicate some of the monospecific clumps of *C. danayanus* colonies. **Fig. 10.** A detail of the moveable plates of the barnacle overgrown with *C. danayanus*. **Fig. 11.** A single cell of *C. danayanus* among dense colony of *Cylindrotheca* sp. attached to the folds in the moveable plates of *P. coriacea*. Scale bars: 10 µm = **Figs 3–5, 7, 11**; 50 µm = **Fig. 2**; 100 µm = **Figs 6, 9 & 10**; 1mm = **Fig. 8**

 **Figures 12–24.** *Craspedostauros danayanus*. **Figs 12–16.** Valve view (light micrographs). Arrows indicate the barely noticeable valve margins. **Figs 17–24.** Scanning electron micrographs. **Fig. 17.**  Detail of the apical part of the valve (external view). Arrowheads indicate the large irregular depression at the fold of the apical silica flap. **Fig. 18.** Frustule with partially detached girdle bands (external view). Arrowheads indicate the large irregular depression at the fold of the apical silica

 flap. **Fig. 19.** Detail of the central part of the valve (external view). **Fig. 20.** Internal valve view. **Fig. 21.** Detail of the central part of the valve (internal view). **Fig. 22.** Cribrate areolae (internal view). **Fig. 23.** Detail of the apical part of the valve (internal view). Arrowheads indicate the asymmetrical thickening extending from the apical part of the raphe-sternum towards the valve margin. **Fig. 24.** Detail of the girdle bands. Scale bars: 10 µm = **Figs 12–16, 18, 20**; 1 µm = **Figs 17, 19, 21–24**

 **Figures 25–40.** *Craspedostauros legouvelloanus*. **Figs 25–30.** Light micrographs. **Figs 25, 26, 28– 30.** Girdle view. **Fig.s 25.** Valve with two girdle bands attached.**, Figs 28 & 29.** Frustules with detached valves. **Figs 26 & 30.** Complete frustules. Arrows indicate the biarcuate valve margin. **Fig. 27.** Valve view. **Figs 31–40.** Scanning electron micrographs. **Fig. 31.** External valve view. Arrows indicate depressions at the apical flap fold. **Fig. 32.** Detail of the apical part of the frustule (external view). **Fig. 33.** Valve with attached girdle bands (girdle view). **Fig. 34.** Detail of the girdle bands (internal view). Arrowheads indicate the internal thickening (septum). **Fig. 35.** Valve with partially detached girdle bands (internal view). **Fig. 36.** Internal valve view. Arrowheads indicate the slight expansion of the stauros on the side corresponding to the external lip-like silica flaps. **Fig. 37.** Detail of the apical part of the valve (internal view). **Fig. 38.** Detail of the central part of the valve (external view). **Figs 39 & 40.** Detail of the central part of the valve (internal view). Arrowheads indicate the hollows in the stauros-adjacent virgae. Scale bars: 10 µm = **Figs 25–31, 33, 35 & 36**; 1 µm = **Figs 32, 34 & 37–39**; 500nm = **Fig. 40** 

 microscopy). Arrows indicate the H-shaped chloroplasts with one lobe pressed against each valve, a feature characteristic of the genus. **Fig. 42.** External valve view (wild population). **Fig. 43.** External valve view (cultured strain). **Fig. 44.** Internal valve view (wild population). **Fig. 45.** Internal valve

**Figures 41–47.** *Craspedostauros legouvelloanus*. **Fig. 41.** Living cells in culture (light

## Page 37 of 113 and 12 and 12 and 13 and 13 and 13 and 13 and 13 and 14 and 15 and 16 Phycology



 **Figures 48–62.** *Craspedostauros macewanii*. **Figs 48–54.** Light micrographs. **Figs 48–51.** Fresh (unpreserved) material. **Figs 48 & 51.** Living cells. **Fig. 48.** Girdle view. **Fig. 51.** Valve view. **Figs 49 & 50.** Damaged cells in girdle view with the cell content (including plastids) spilling beyond the cell wall. **Figs 49.** Arrow indicates the straight valve margin. **Figs 52–54.** Cleaned material. Detached valves in valve view. Arrows indicate the distinct valve face-mantle junction. **Figs 55–62.**  Scanning electron micrographs. **Fig. 55.** External valve view. **Fig. 56.** Detail of the apical part (external valve view). **Fig. 57.** Detail of the central area (external valve view). **Fig. 58.** Detail of the apical part (external girdle view). **Fig. 59.** Internal valve view and partially detached valvocopula. **Fig. 60.** Detail of the apical part (internal valve view). Arrowheads indicate several small areolae present at the end of the curved thickening. **Fig. 61.** Detail of the central area (internal valve view). **Fig. 62.** Detail of the valvocopula (internal view).

Scale bars: 10 µm = **Figs 48–55 & 59**; 1 µm = **Figs 56–58 & 60–62**

 **Figures 63–74.** *Craspedostauros alatus* (Adriatic population). **Figs 63–68.** Light micrographs. **Figs 63, 66 & 67.** Valve view. **Fig. 63.** Broken frustule with both valves lying in valve view. **Fig. 64.**  Single valve with attached girdle bands. **Figs 65 & 68.** Girdle view. Arrows indicate the clear valve face-mantle junction. **Figs 69–74.** Scanning electron micrographs. **Fig. 69.** Frustule with partially detached girdles bands (external view). **Fig. 70.** Detail of the apical part of the frustule with the winged-liked silica flaps, a feature typical of the species (external view). **Fig. 71.** Frustule with partially detached girdles bands (external girdle view). **Fig. 72.** Internal valve view. **Figs 73 & 74.**

 Detail of the central part of the valve (internal view). Scale bars: 10 µm = **Figs 63–69, 71 & 72**; 1 µm = **Figs 70 & 73**; 500 nm = **Fig. 74**

 **Figure 75.** Maximum likelihood (ML) phylogram based on the 3-gene dataset (nuclear-encoded ribosomal SSU, chloroplast encoded rbcL, psbC markers). For clarity, only the clade of raphid diatoms containing *Staurotropis*, *Craspedostauros*, and *Achnanthes* is presented in the figure. The ML tree presenting the complete taxon sampling can be viewed in the Supplementary Figure S1. 

## **Supplementary Figure S1**

 Maximum likelihood tree based on the 3-gene dataset (nuclear-encoded ribosomal SSU, chloroplast-encoded rbcL, psbC markers) with bootstrap values from 1000 pseudoreplicates over the corresponding nodes. The araphid pennate taxon outgroup *Asterionellopsis socialis* was used as the outgroup.



Sampling locations where Craspedostauros danayanus (1), C. legouvelloanus (2), C. macewanii (3), and C. alatus (4) were found.

123x160mm (600 x 600 DPI)



Craspedostauros danayanus. Fig. 2. Living cells of C. danayanus and Cylindrotheca sp. attached to the leatherback skin scutes (light microscopy). Fig. 3. Stained colony of C. danayanus and associated bacteria on the leatherback skin scutes. Fig. 4. Valve view of a living cell (cultured strain). Fig. 5. Girdle view of a living cell (cultured strain). Figs 6–11. Scanning electron micrographs of C. danayanus attached to its original substratum. Fig. 6. Monospecific colony growing among the flaking skin of leatherback (dorsal side of the hind flipper). Fig. 7. Extremely delicate and fragile cells of C. danayanus attached to the leatherback skin (dorsal side of the hind flipper). Fig. 8. An overview of the leatherback-associated barnacle, Platylepas coriacea, colonized by C. danayanus. Fig. 9. A detail of the external part of the barnacle with a sheath of host sea turtle tissue overgrown with C. danayanus. Arrows indicate some of the monospecific clumps of C. danayanus colonies. Fig. 10. A detail of the moveable plates of the barnacle overgrown with C. danayanus. Fig. 11. A single cell of C. danayanus among dense colony of Cylindrotheca sp. attached to the folds in the moveable plates of P. coriacea. Scale bars:  $10 \mu m =$  Figs 3-5, 7, 11; 50  $\mu m =$  Fig. 2; 100  $\mu m =$  Figs 6, 9 & 10;  $1mm = Fig. 8$ 



Craspedostauros danayanus. Figs 12–16. Valve view (light micrographs). Arrows indicate the barely noticeable valve margins. Figs 17–24. Scanning electron micrographs. Fig. 17. Detail of the apical part of the valve (external view). Arrowheads indicate the large irregular depression at the fold of the apical silica flap. Fig. 18. Frustule with partially detached girdle bands (external view). Arrowheads indicate the large irregular depression at the fold of the apical silica flap. Fig. 19. Detail of the central part of the valve (external view). Fig. 20. Internal valve view. Fig. 21. Detail of the central part of the valve (internal view). Fig. 22. Cribrate areolae (internal view). Fig. 23. Detail of the apical part of the valve (internal view). Arrowheads indicate the asymmetrical thickening extending from the apical part of the raphe-sternum towards the valve margin. Fig. 24. Detail of the girdle bands. Scale bars: 10 µm = Figs 12–16, 18, 20; 1 µm = Figs 17, 19, 21–24



Craspedostauros legouvelloanus. Figs 25–30. Light micrographs. Figs 25, 26, 28–30. Girdle view. Fig. 25. Valve with two girdle bands attached. Figs 28 & 29. Frustules with detached valves. Figs 26 & 30. Complete frustules. Arrows indicate the biarcuate valve margin. Fig. 27. Valve view. Figs 31–40. Scanning electron micrographs. Fig. 31. External valve view. Arrows indicate depressions at the apical flap fold. Fig. 32. Detail of the apical part of the frustule (external view). Fig. 33. Valve with attached girdle bands (girdle view). Fig. 34. Detail of the girdle bands (internal view). Arrowheads indicate the internal thickening (septum). Fig. 35. Valve with partially detached girdle bands (internal view). Fig. 36. Internal valve view. Arrowheads indicate the slight expansion of the stauros on the side corresponding to the external lip-like silica flaps. Fig. 37. Detail of the apical part of the valve (internal view). Fig. 38. Detail of the central part of the valve (external view). Figs 39 & 40. Detail of the central part of the valve (internal view). Arrowheads indicate the hollows in the stauros-adjacent virgae. Scale bars:  $10 \mu m =$  Figs 25-31, 33, 35 & 36; 1  $\mu m =$  Figs 32, 34 & 37-39;  $500$ nm = Fig. 40



Craspedostauros legouvelloanus. Fig. 41. Living cells in culture (light microscopy). Arrows indicate the Hshaped chloroplasts with one lobe pressed against each valve, a feature characteristic of the genus. Fig. 42. External valve view (wild population). Fig. 43. External valve view (cultured strain). Fig. 44. Internal valve view (wild population). Fig. 45. Internal valve view (cultured strain). Fig. 46. Detail of a girdle band showing internal thickening (septum) with perforations. Fig. 47. A single girdle band (external and internal view). Scale bars: 10  $\mu$ m = Figs 41-45; 1  $\mu$ m = Figs 46 & 47



Craspedostauros macewanii. Figs 48–54. Light micrographs. Figs 48–51. Fresh (unpreserved) material. Figs 48 & 51. Living cells. Fig. 48. Girdle view. Fig. 51. Valve view. Figs 49 & 50. Damaged cells in girdle view with the cell content (including plastids) spilling beyond the cell wall. Figs 49. Arrow indicates the straight valve margin. Figs 52–54. Cleaned material. Detached valves in valve view. Arrows indicate the distinct valve face-mantle junction. Figs 55–62. Scanning electron micrographs. Fig. 55. External valve view. Fig. 56. Detail of the apical part (external valve view). Fig. 57. Detail of the central area (external valve view). Fig. 58. Detail of the apical part (external girdle view). Fig. 59. Internal valve view and partially detached valvocopula. Fig. 60. Detail of the apical part (internal valve view). Arrowheads indicate several small areolae present at the end of the curved thickening. Fig. 61. Detail of the central area (internal valve view).

Fig. 62. Detail of the valvocopula (internal view). Scale bars: 10  $\mu$ m = Figs 48-55 & 59; 1  $\mu$ m = Figs 56-58 & 60-62



Craspedostauros alatus (Adriatic population). Figs 63–68. Light micrographs. Figs 63, 66 & 67. Valve view. Fig. 63. Broken frustule with both valves lying in valve view. Fig. 64. Single valve with attached girdle bands. Figs 65 & 68. Girdle view. Arrows indicate the clear valve face-mantle junction. Figs 69–74. Scanning electron micrographs. Fig. 69. Frustule with partially detached girdles bands (external view). Fig. 70. Detail of the apical part of the frustule with the winged-liked silica flaps, a feature typical of the species (external view). Fig. 71. Frustule with partially detached girdles bands (external girdle view). Fig. 72. Internal valve view. Figs 73 & 74. Detail of the central part of the valve (internal view). Scale bars: 10 µm = Figs 63–69, 71 & 72; 1 µm = Figs 70 & 73; 500 nm = Fig. 74



Maximum likelihood (ML) phylogram based on the 3-gene dataset (nuclear-encoded ribosomal SSU, chloroplast encoded rbcL, psbC markers). For clarity, only the clade of raphid diatoms containing Staurotropis, Craspedostauros, and Achnanthes is presented in the figure. The ML tree presenting the complete taxon sampling can be viewed in the Supplementary Figure S1.

170x66mm (300 x 300 DPI)

## **Table 1**

Comparison of *Craspedostauros alatus*, *C. danayanus, C. legouvelloanus*, and *C. macewanii* with several morphologically similar *Craspedostauros* taxa (after Cox 1999, Ashworth et al. 2017, and Majewska et al. 2018).



\* values and descriptions given in brackets refer to the Adriatic populations



0.08

Table S1. Taxa, strain voucher ID and GenBank accession numbers for strains used in the DNA sequence data phylogenetic analysis. Collection site for sample of original strain isolation is also included (where known); in the case of cultures from public collections, the culture ID is provided in this column (UTEX = UTEX Culture Collection of Algae; NCMA = National Center for Marine Algae and Microbiota; CSIRO = Australian National Algae Culture Collection; MCC-NIES = Microbial Culture Collection at National Institute for Environmental Studies). Ingroup taxa (raphid pennates) provided first in the table; outgroup taxa ("araphid pennates") follow after table break. Taxa are listed alphabetically. If species unknown, authority for genus is listed.
























































Running title: Sea turtle-associated *Craspedostauros*

## **ABSTRACT**

 Despite recent advances in the research on sea turtle-associated diatoms, some of the key aspects of the diatom-sea turtle relationship, including compositional and functional features of the epizoic diatom community, remain understudied and poorly understood. The current paper focuses on four species belonging to the primarily marine diatom genus *Craspedostauros* that were observed growing attached to numerous sea turtles and sea turtle-associated barnacles from Croatia and South Africa. Three of the examined taxa, *C. danayanus* sp. nov., *C. legouvelloanus* sp. nov., and *C. macewanii* sp. nov. represent novel species and are described based on morphological and, whenever possible, molecular characteristics. The new taxa exhibit characters not yet observed in other members of the genus, such as the presence of more than two rows of cribrate areolae on the girdle bands, shallow perforated septa, and a complete reduction of the stauros. In addition, *C. alatus*, recently described from museum sea turtle specimens, is reported for the first time from loggerheads rescued in Europe. A 3-gene phylogenetic analysis including DNA sequence data for three sea turtle-associated *Craspedostauros* species and other marine and epizoic diatom taxa indicated that *Craspedostauros* is monophyletic and sister to *Achnanthes*. This study, being based on a large number of samples and animal specimens analysed and using different preservation and processing methods, provides some new insights into the genus ecology and biogeography and sheds more light on the level of intimacy and permanency in the host-epibiont interaction within the epizoic *Craspedostauros* species.

 **Key index words:** *Craspedostauros*, barnacle, *Chelonibia*, epizoic diatom, leatherback, loggerhead, phylogeny, *Platylepas*, sea turtle

 **Abbreviations:** BS, bootstrap support; CRW, Comparative RNA Web; LM, light microscopy; ML, maximum likelihood; SEM, scanning electron microscopy; SSU, small subunit

#### **INTRODUCTION**

 As indicated by several studies, diatom communities inhabiting both the skin and the carapace of marine turtles are composed largely of species not observed on other biotic or abiotic substrata (Frankovich et al. 2015, 2016, Majewska et al. 2015a, 2015b, 2017a, 2017b, Robinson et al. 2016, Azari et al. 2020). These observations further suggest a certain level of host-specific evolutionary adaptations used by diatoms. Although intimate relationships between animals and microbes are common and extensively studied, reports of truly epizoic microalgae are generally rare (Ezenwa et al. 2012, Redford et al. 2012, Apprill 2017). Perhaps due to the fact that ubiquitous photosynthetic organisms, such as diatoms, are not immediately perceived as an essential element of any vertebrate microbiome, these new findings are particularly noteworthy. Based on their high frequency of occurrence and high relative abundances recorded from various sea turtle species and geographical regions, as well as lack of records from other types of substrata, several of the newly described sea turtle-associated diatom taxa are currently believed to be strictly epizoic or even sea turtle-specific. While this may be true, many other diatoms present in the sea turtle samples are likely opportunistic species that attached to biofilm in the later stages of its development (Majewska et al. 2015b, 2017b, 2019a,b). Although opportunistic taxa often dominate specific epizoic habitats in terms of the species number, they rarely reach high relative abundance, which may suggest their lack of some key functional adaptations to the epizoic lifestyle.

The present study focuses on the sea turtle-associated species belonging to the diatom genus

*Craspedostauros* E.J.Cox. At present, the genus comprises ten validly described species including

one, *C. alatus* Majewska et Ashworth, described from museum specimens of sea turtles (Cox 1999,

Sabbe et al. 2003, Van de Vijver et al. 2012, Ashworth et al. 2017, Majewska et al. 2018).

*Craspedostauros* is a predominantly marine genus, although *C. laevissimus* (W. et G.S. West)

Sabbe is described as "a widespread endemic species restricted to the Antarctic Continent" and may

#### Page 81 of 113 and 12 and 12 and 13 and 1

 be of brackish or freshwater origin (Sabbe et al. 2003, Van de Vijver et al. 2012). Most of the *Craspedostauros* members share the typical of the genus morphological characters such as cribrate areolae, numerous doubly-perforated girdle bands, two fore and aft chloroplasts, and a usually narrow stauros. Nevertheless, the latter is reduced or strongly reduced in two species: *C. alyoubii* J.Sabir et Ashworth and *C. paradoxus*\* Ashworth et Lobban. Molecular phylogenetic analysis indicated that the genus is closely related to *Achnanthes* Bory and *Staurotropis* Paddock (Ashworth et al. 2017). Both taxa, as well as another marine genus *Druehlago* Lobban et Ashworth, which has yet to be characterized molecularly, share several morphological similarities with *Craspedostauros* (Cox 1999, Ashworth et al. 2017). For example, all the above-mentioned taxa possess valves and girdle bands perforated by cribrate areolae. Moreover, *Craspedostauros* and *Druehlago* share the general frustule morphology, including frustules with central constriction (Ashworth et al. 2017), whereas the fore and aft arrangement of chloroplasts, typical of *Craspedostauros,* can be observed in several *Achnanthes* species (Cox 1999).Three novel species, *C. danayanus* Majewska et Ashworth sp. nov., *C. legouvelloanus* Majewska et Bosak sp. nov., and *C. macewanii* Majewska et Ashworth sp. nov., were found in the course of the ongoing survey on sea turtle-associated diatoms and are described in the current paper. Moreover, a small population of *C. alatus* is for the first time reported from Europe. A large number of samples analysed and different preservation and processing techniques applied allowed us to document the ultrastructure of the frustule and, whenever possible, the morphology of the plastids as well as the colony type and attachment mode of the cells. These observations were supplemented by a 3-gene phylogenetic analysis including DNA sequence data for three sea turtle-associated *Craspedostauros* species and other marine and epizoic diatom taxa.

 \* the specific epithet in *Craspedostauros paradoxa* should be changed to '*paradoxus*' following the recommendations of the International Code of Nomenclature for algae, fungi, and plants (Articles 23.5 & 62; Turland et al. 2018).

### **MATERIALS AND METHODS**

### *Material collection and preservation*

 Diatom samples were collected from captive and wild sea turtles from Croatia and South Africa. All biofilm samples from carapace and skin were taken using single-use sterile toothbrushes according to the sampling protocols suitable for diatom culturing and standard morphology-based diatom analysis proposed by Pinou et al. (2019). In Croatia, 76 (skin and carapace) samples were collected from 38 loggerhead sea turtles *Caretta caretta* L. rescued and rehabilitated at the Marine Turtle Rescue Centre in Aquarium Pula between 2016 and 2019, on the day of or shortly after their arrival at the facility. In South Africa, 196 (skin and carapace) biofilm samples were collected from 78 loggerheads and 20 leatherbacks *Dermochelys coriacea* Vandelli nesting in Kosi Bay (Indian Ocean) over two nesting seasons, in 2017/2018 and 2018/2019. In addition, 6-mm skin biopsy punches were taken from either front or rear flippers of 30 loggerheads and six leatherbacks and preserved in 4 % formaldehyde solution in seawater immediately after collection. Samples of sea turtle-associated barnacles *Chelonibia testudinaria* L. from 100+ loggerheads and *Platylepas coriacea* Monroe et Limpus from 15 leatherbacks were taken using a plastic paint scraper or a blunt knife during four nesting seasons, in 2015/2016, 2016/2017, 2017/2018, and 2018/2019. Barnacle samples comprised of more than one specimen, were divided into two parts and either frozen (- 20°C) or fixed with 4 % formaldehyde solution in seawater. Single-specimen barnacle samples were frozen (-20°C). Furthermore, skin and carapace samples were collected from seven sea turtles (three loggerheads, three green turtles *Chelonia mydas* L., and one hawksbill *Eretmochelys imbricata* L.) resident at the uShaka Sea World in Durban on 28 June 2019.

#### Page 83 of 113 and 12 and 13 and 13 and 13 and 13 and 14 and 15 and 16 and 16 Phycology



*Material processing and microscopy*

 Diatoms were detached from the frozen barnacles using a Transsonic T310 (Elma, Singen, Germany) ultrasound bath as described in Majewska et al. (2019b). Diatom biofilm from the sea turtle skin, carapace, and barnacles was cleaned from organic matter using either a rapid digestion 150 with a mixture of concentrated  $HNO_3$  and  $H_2SO_4$  (at a ratio of 2:1) according to the method proposed by von Stosch (South African and Croatian samples; Hasle and Syvertsen 1997) or heated  $37\%$  H<sub>2</sub>O<sub>2</sub> with addition of KMnO<sub>4</sub> (Croatian culture strain; van der Werff 1953). Cleaned material was mounted on slides using Naphrax (Brunel Microscopes Ltd, Chippenham, UK; Croatian samples) and Pleurax prepared according to the method proposed by von Stosch (1974; South African samples). The slides were examined using a Nikon Eclipse 80i light microscope with Differential Interference Contrast (DIC) and a Nikon DS-Fi1 5MP digital camera (Nikon Instruments Inc., Melville, NY; South African samples) as well as a Zeiss Axio Imager A2 with DIC and an Axiocam 305 digital camera (Carl Zeiss, Jena, Germany; Croatian samples). In addition, fresh material containing living diatoms attached to the sea turtle scutes and skin flakes was stained with blue writing ink (Scheaffer ®) to reveal the colonies of the diatom-associated bacteria.

 For scanning electron microscopy (SEM), the oxidized suspension was filtered through 1-µm or 1.2-µm Isopore™ (Merck Millipore, Darmstadt, Germany) or 3-µm Nucleopore (Nucleopore, Pleasanton, CA, USA) polycarbonate membrane filters. Formalin-preserved skin and barnacle samples were dehydrated in an alcohol series (30%, 50%, 60%, 70%, 80%, 90%, 95%, 99.9%) followed by critical point-drying in an E3100 Critical Point Dryer (Microscience Division, Watford,

#### Journal of Phycology **Page 84 of 113**



### *Culturing*

 Living diatoms from the fresh material (unpreserved samples containing sea turtle biofilm and filtered seawater; Pinou et al. 2019) were isolated using a glass pipette with a tip pulled and thinned over a flame into 16x100 mm glass culture tubes (South African strains) or plastic culture flasks (Croatian strains) filled with 34 PSU (South African strains) or 38 PSU (Croatian strains) f/2 growth medium (Guillard 1975). Strains were lit by natural light from a south-facing window (South African strains) or white fluorescent light with a photoperiod of 12h (Croatian strains) and maintained at a temperature of 20*–*24°C. The well-growing cultures were divided into two parts, one of which was used for DNA extraction. The remaining part was cleaned with a mixture of 30%  $H_2O_2$  and 70% HNO<sub>3</sub> and rinsed with distilled water until the near-neutral pH of the fluid phase was 191 reached. Croatian strain (PMFTB0003) was cleaned using saturated KMnO<sub>4</sub> solution and ca. 30%

#### Page 85 of 113 and 12 and 13 and 13 and 13 and 13 and 13 and 13 and 14 and 15 and 16 Phycology

 HCl following a slightly modified protocol proposed by Simonsen (1974). Permanent microscopy slides and SEM stubs were prepared as described above.

### *DNA preparation and phylogenetic analysis*

 The cultures were harvested as cell pellets using an Eppendorf 5415C centrifuge (Eppendorf North America, Hauppauge, NY, USA) for 10 minutes at 8000 rpm. The QIAGEN DNeasy Plant Mini Kit (QIAGEN Sciences, Valencia, California, USA) was used for DNA extraction following the manufacturer's protocol, with the addition of an initial cell disruption by 1.0 mm glass beads in a Mini-Beadbeater (Biospec Products, Inc, Bartlesville, OK, USA) for 45 sec. PCR-based DNA amplification and di-deoxy Sanger sequencing of small-subunit nuclear rRNA and the chloroplast-encoded rbcL and psbC markers followed Theriot et al. (2010).

 Phylogenetic analysis of the DNA sequence data was conducted using a three-gene dataset: nuclear- encoded small subunit (SSU) rRNA, and plastid-encoded *rbc*L and *psb*C. Alignment of the SSU sequences, accounting for secondary structure, was done using the SSUalign program (Nawrocki et al. 2009), with the covariance model based on the 10 diatoms included with the program download, plus 23 additional diatoms from the CRW website (Cannone et al. 2002). Post alignment, SSU sequences were concatenated to the chloroplast sequences into a single matrix (Supplementary Table S1). Eight separate partitions were created for the data (SSU paired and unpaired sites, plus the first, second and third codon positions of each of *rbc*L and *psb*C). This dataset and partitioning scheme were run under maximum likelihood (ML) using RAxML ver. 8.2.7 (Stamatakis 2014) 212 compiled as the pthread-AVX version on an Intel i7 based processor, using the GTR+G model. Twenty-five replicates, each with 500 rapid BS replicates, were run with ML optimizations. Bootstrap support was assessed using the BS replicates from the run with the optimal ML score.

#### **RESULTS**

*Morphological observations*

## *Craspedostauros danayanus* **Majewska & Ashworth sp. nov. (Figs 2–24)**

Cells with two fore and aft H-shaped chloroplasts (Figs 2*–*5). Frustules extremely delicate and very

lightly silicified (Figs 6*–*16). In girdle view, frustules rectangular, moderately constricted at the

centre (Figs 5, 7 & 11). Valves narrow, linear, very slightly constricted in the valve middle, with

bluntly rounded apices (Figs 4, 12*–*16).

# *Light microscopy (Figs 12–16):*

 Valve dimensions (*n* = 30): length 28*–*61 μm, width 2–2.5 μm, length/width ratio: 14–30.5. In 226 cleaned (acid-digested) material, partially dissolved valve margins barely noticeable (Figs 14  $&$  15, arrows), intact frustules absent. Striae indiscernible (Figs 12*–*16). Raphe-sternum thickened, clearly visible (Figs 12*–*16). Thickenings at both central and terminal raphe endings (Figs 12*–*16).

# *Scanning electron microscopy (Figs 17–24):*

 Externally: In cleaned material, valve face appearing flat, with very shallow mantle and straight margin (Figs 17 & 18). Striae uniseriate, 49*–*51 in 10 μm, parallel, becoming radiate towards the 233 apices, alternate or opposite, composed of up to eight areolae (Figs 17 & 18). Areolae largely similar in size, becoming somewhat smaller around the central area, squarish to roundish, externally occluded by cribra (Figs 17–19). Each cribrum perforated by 2–8 pores (Fig. 17). Axial area narrow (Figs 17 & 18). Raphe-sternum not raised (Figs 17–19). Raphe branches straight (Fig. 18). Central area large, symmetrical, amygdaliform (Figs 18 & 19). Central raphe endings straight, elongated, slightly expanded (Figs 18 & 19). Terminal raphe endings disappearing under somewhat triangular

#### Page 87 of 113 and 113 and 12 Journal of Phycology



 roundish or elongated areolae, ca. 50*–*60 in 10 μm (Figs 18, 23 & 24). Areolae occluded externally by cribra (Figs 23 & 24).

### **Taxonomic remarks**

 *Craspedostauros danayanus* is most similar to *C. paradoxus*, sharing the general valve outline and lacking the stauros. However, *C. danayanus* differs from the latter in being distinctly smaller (28*–*

- 61 μm vs 80*–*85 μm) and more slender (2*–*2.5 μm vs 6.5*–*9 μm), possessing a higher stria density
- (49*–*51 vs 36*–*40), and lacking the lip-like silica flaps (externally) and the central knob (internally)

present in *C. paradoxus* (Table 1).



deposited in the South African Diatom Collection housed by North-West University,

Potchefstroom, South Africa.

266 TYPE LOCALITY: Mabibi Beach, Elephant Coast, South Africa (27° 21′ 30" S, 32° 44′ 20" E).

Collected from the barnacle *Platylepas coriacea* growing on the egg-lying leatherback sea turtle

(tag numbers: ZA0019A, ZA1824E) by R. Majewska, 7 December 2018.

ETYMOLOGY: The epithet honours Danay A. Stoppel (North-West University, Potchefstroom,

South Africa), who made the first observations of the new taxon, in recognition of her contribution

to the sea turtle diatom project in South Africa.

ECOLOGY: Epizoic on carapaces of adult leatherback sea turtles and on leatherback-associated

barnacles *Platylepas coriacea* growing on adult leatherbacks from Kosi Bay (South Africa).

Attaching to the animal surface through one end of the valve, motile in culture.

 The taxon was found in twelve leatherback skin samples (out of 20 examined) and in all *P. coriacea* 276 samples examined ( $n = 15$ ) reaching relative abundances of 35% (skin samples) and 79% (barnacle samples). It was found in neither loggerhead nor loggerhead-associated barnacle samples from the same location (Kosi Bay, South Africa). Leatherback skin samples containing *C. danayanus* were dominated by *Navicula* spp., *Tursiocola* sp., and *Poulinea* spp. The new taxon was dominant in most of the *P. coriacea* samples along with *Cylindrotheca* sp. Both taxa colonised various anatomical parts of the barnacle showing preference for rough surfaces and cavities. The extremely lightly silicified frustules may be an adaptation to the pelagic lifestyle of the host, as the open ocean waters contain significantly lower concentrations of dissolved silica than coastal habitats (Tréguer et al. 1995).

*Craspedostauros legouvelloanus* **Majewska & Bosak sp. nov. (Figs 25–47)**

# *Light microscopy (Figs 25–30):*

 Intact frustules lying almost always in girdle view (due to large cell depth/valve width ratio), slightly constricted in the middle (Figs 25, 26, 28*–*30), with several girdle bands (Figs 26, 28 & 30). Valve margin expanded at the centre (Figs 25, 28 & 30). Frustules lightly silicified and delicate. Valves narrow, linear to linear-lanceolate, slightly constricted at the central area, with bluntly rounded apices (Fig. 27). Valve dimensions (*n* = 30): length 18*–*34 μm, width 3–5 μm, length/width ratio: 5.6–9.4. Striae indiscernible (Figs 25*–*30). Stauros narrow (Figs 25, 27*–*30), widening towards the biarcuate valve margins (Fig. 30, arrows). Raphe-sternum clearly visible (Figs 25*–*30). Raphe 295 straight, biarcuate in girdle view (Figs 25, 26, 28 & 30).

### *Scanning electron microscopy (Figs 31–40):*

 Externally: Valves somewhat convex, with no clear valve face-mantle junction (Figs 31–33). Valve margin clearly expanded at the centre beyond the stauros (Fig. 33). Striae uniseriate, 46*–*49 in 10 μm, parallel throughout the valve centre, becoming convergent near the apices, alternate or opposite, composed of up to 13 areolae (Figs 31, 32 & 38). Areolae similar in size throughout the entire valve, squarish, externally occluded by cribra (Figs 31–33 & 38). Each cribrum perforated by 4 pores (Figs 31–33 & 38). Axial area very narrow (Figs 31 & 32). Raphe-sternum very slightly raised (Figs 31–33). Raphe branches more or less straight (Fig. 31). Central area forming a narrow rectangular fascia (Figs 31 & 38). Central raphe endings covered entirely by rimmed lip-like silica 306 flaps extending from one side of the axial area (Figs 31  $\&$  38). At the apices, axial area expanding into somewhat triangular silica flaps covering the terminal raphe endings giving the impression of unilaterally bent terminal raphe fissures (Figs 31–33). An oval or irregular depression present at the apical flap fold (Fig. 31, arrows). Shortened stria composed of regular areolae and simple puncta radiating around the apices beyond the terminal raphe endings (Figs 31–33).

#### Journal of Phycology **Page 90 of 113**

 Internally: Raphe slit opening laterally onto the uniformly thick and clearly raised raphe-sternum (Figs 35 & 36). Stauros raised, very narrow, broadening abruptly at the mantle expansion and merging with the pore-free area at the valve margin (Figs 36 & 39), slightly more expanded on the side corresponding to the external lip-like silica flaps (Figs 36, arrowheads, 39 & 40). Central raphe endings straight or slightly unilaterally bent, terminating onto weakly developed, elongated and flattened helictoglossae (Figs 35, 36, 39 & 40). A blunt cylindrical knob with a small central cavity present between the raphe endings (Figs 35, 36, 39 & 40). Areolae externally occluded by cribra, 318 appearing sunken, especially close to the stauros (Figs  $39 \& 40$ ). Stauros-adjacent virgae appearing hollow, suggesting a more complex valve structure in that area (Fig. 39, arrowheads). Terminal raphe endings positioned somewhat laterally on the raphe-sternum, terminating onto prominent helictoglossae. At the apices, raphe-sternum expanded laterally towards the valve margin, merged 322 with pore-free area corresponding to the external apical silica flaps (Figs 36 & 37). Cingulum composed of numerous (12+) open copulae, bearing two rows of typically squarish or elongated areolae, ca. 50*–*60 in 10 μm (Figs 32–35). Areolae occluded externally by cribra with 4*–* 12 pores per cribrum (Figs 32–35). Valvocopula curved, distinctly narrower and pore-free beside

 the stauros (Fig. 33, arrowheads). An internal ridge perforated by puncta, resembling a reduced septum, present in each copula except for valvocopula (Figs 33–35, arrowheads).

### *Adriatic population (Figs 41–47)*

 Specimens resembling *C. leguovelloanus* were found on the carapace of six loggerhead sea turtles sampled on the Croatian coast of the Adriatic Sea. Most of the morphological features observed in the Adriatic population (Figs 41–47) agreed well with those found in *C. legouvelloanus*. The cells possessed two fore and aft H-shaped chloroplasts (Fig. 41, arrows) observed previously in other *Craspedostauros* species (Cox 1999, Ashworth et al. 2017, Majewska et al. 2018). The specimens were slightly longer (23*–*39 μm) and wider (3.5*–*6 μm, length/width ratio: 5.2*–*7.8, *n* = 25) than

#### Page 91 of 113 and 12 and 12 and 13 and 14 and 15 and 16 Phycology

 those from the South African population and their stria density was lower (40*–*44 in 10 μm vs. 46*–* 49 in 10 μm; Table 1). In general, the frustules showed a relatively high degree of irregularity in the areolae structure and the size and shape of stauros, axial area, and facia (Figs 42*–*45).

## **Taxonomic remarks**

 Currently, *C. legouvelloanus* is the only *Craspedostauros* species with septate girdle bands. Valves of this species differ from those of all known stauros-bearing *Craspedostauros* species in possessing a very high stria density (above 40 in 10 μm). Although a similarly high or higher stria density was observed in *C. alyoubii* (~40 in 10 μm) and *C. danayanus* (49*–*51 in 10 μm), the two species are larger (83*–*105 μm and 28*–*61 μm) than *C. legouvelloanus* (18*–*34 [39] μm) and their general morphology differs remarkably from that of the new taxon in, for example, possessing a reduced or strongly reduced stauros (Table 1). Several of the characters of *C. legouvelloanus*, such as largely uniform valve areolae with four pores per cribrum and internal central knob, agree with the description of *C. australis* E.J.Cox (Cox 1999)*.* However, the new species can be easily distinguished from the latter by its clearly centrally expanded valve margin and well-developed lip- like silica flaps externally covering the central raphe endings absent in *C. australis* (Table 1). Although wild specimens belonging to the Adriatic population of *C. legouvelloanus* exhibited numerous irregularities in the shape and size of taxonomically important characters such as areolae, striae, stauros, and central area, we were unable to indicate and unambiguously describe features that would distinguish them from the type population. High morphological plasticity and polymorphy in diatoms have been reported from both epizoic and non-epizoic habitats (Cox 2011, De Martino et al. 2011, Urbánková et al. 2016, Riaux-Gobin et al. 2014, 2017, Edlund and Burge 2019), and it is conceivable that the morphological differences observed between the two

populations could be induced by environmental triggers, such as differences in salinity or nutrient

concentrations (Schultz 1971, Czarnecki 1987, 1994, De Martino et al. 2011). Unfortunately, the

 Croatian strain PMFTB0003 (Figs 41, 43, 45 & 46) isolated from the sample TB13 did not survive and the DNA material could not be obtained at the time of this study. Therefore, in the light of the current lack of any additional information about the phylogenetic relationships between the two populations, they should be considered conspecific until otherwise proven.

 HOLOTYPE: Permanent slide SANDC-ST003 and unmounted material (prepared from sample ZA0762D/ZA0763D) deposited in the South African Diatom Collection housed by North-West University, Potchefstroom, South Africa.

PARATYPE: Permanent slide HRNDC000150 and unmounted material (TB13) deposited in the

Croatian National Diatom Collection housed by Faculty of Science, University of Zagreb, Croatia.

ISOTYPES: Permanent slides BR-XXXX and BR-XXXX deposited in the BR-collection housed by

Meise Botanic Garden, Meise, Belgium.

373 TYPE LOCALITY: Kosi Bay, South Africa (26° 59' 39" S, 32° 51' 60" E). Collected from the

carapace of the egg-lying loggerhead sea turtle (tag numbers: ZA0762D, ZA0763D) by R.

Majewska, 15 December 2017 (holotype).

376 Marine Turtle Rescue Centre, Pula, Croatia (44°50′ 07" N, 13°49′ 58" E). Collected from a semi- adult female loggerhead *Caretta caretta* named 'Mimi' by K. Gobić Medica, 28 May 2019 (paratype).

 ETYMOLOGY: The epithet honours Dr Diane Z. M. Le Gouvello du Timat (Nelson Mandela University, Port Elizabeth, South Africa), who assisted during the type material collection, in recognition of her invaluable help and on-going support to the sea turtle diatom project and sea turtle research in South Africa.

ECOLOGY: Epizoic on carapaces and skin of adult loggerhead sea turtles and on loggerhead-

associated barnacles *Chelonibia testudinaria* growing on adult loggerheads from Kosi Bay (South



 Although the taxon was present in numerous samples, its relative abundance rarely exceeded 4% of the total diatom number. Samples with *C. legouvelloanus* from both locations were each time dominated by *Poulinea* spp., *Berkeleya* spp., *Halamphora* spp., and *Nitzschia* spp., with addition of *Achnanthes elongata* Majewska et Van de Vijver, *Cyclophora tenuis* Castracane, *Proschkinia* spp., *Navicula* spp., *Licmophora* spp., and *Haslea* spp.

## *Craspedostauros macewanii* **Majewska & Ashworth sp. nov. (Figs 48–62)**

## *Light microscopy (Figs 48–54):*

 Cells with two fore and aft H-shaped chloroplasts (Figs 48 & 51). Frustules delicate and lightly silicified (Figs 48*–*54). In girdle view, frustules rectangular, moderately to strongly constricted at the centre (Figs 48*–*50). Cingulum composed of several girdle bands (Figs 49*–*50). Valves narrow, linear to linear-lanceolate, slightly constricted at the central area, with bluntly rounded apices (Figs 51*–*54). Valve margin straight (Fig. 49, arrow). Valve dimensions (*n* = 20): length 26–51 μm (up to 65 μm in culture), width 4.5–5.5 μm (up to 6 μm in culture), length/width ratio: 5.4–11.3. Valve face-mantle junction visible on each side of the raphe (Figs 52–54, arrows). Striae barely discernible, 28–31 in 10 μm (Figs 52*–*54). Central area narrow, bow tie-shaped (Figs 52*–*54). Raphe-sternum thickened (Figs 52*–*54). Raphe straight (Fig. 54) with thickenings at the terminal raphe endings (Figs 52*–*54).

## *Scanning electron microscopy (Figs 55–62):*

Externally: Valves slightly concave at the centre, with distinct valve face-mantle junction marked

by a narrow pore-free area (Figs 55 & 57). Valve face flat (Fig. 55). Mantle very deep (Fig. 55).

#### Journal of Phycology **Page 94 of 113**

409 Valve margin straight, with narrow pore-free area at the mantle edge (Figs 56  $&$  57). Striae uniseriate, parallel through most of the valve, becoming convergent near the apices, alternate or opposite, composed of up to 21 areolae (2–8 on the valve face and up to 13 on the mantle; Figs 55– 58). Areolae similar in size, squarish, externally occluded by cribra (Figs 56–58). Areolae bordering the narrow axial area usually only slightly larger and somewhat irregular in shape (Figs 56–58). Each cribrum perforated by highly variable number of pores (up to 13+; Figs 56–58). Raphe branches more or less straight (Fig. 55). Central area in the form of a narrow bow tie-shaped fascia (Figs 55 & 57). Central raphe endings covered by small lip-like silica flaps extending from one side 417 of the axial area (Figs 55  $\&$  57). Apices pore-free (Figs 55, 56  $\&$  58). Terminal raphe endings covered by triangular silica flaps giving the impression of unilaterally bent terminal raphe fissures (Figs 55, 56 & 58). An oval or irregular depression (Figs 55, arrowhead, 56 & 58) with several small areolae (Figs 56 & 58, arrowheads) present at the apical flap fold. Shortened striae composed of a single areola (occasionally with additional puncta) radiating around the apices beyond the terminal raphe endings (Figs 56 & 58).

 Internally: Raphe slit opening more or less centrally onto the uniformly thick raphe-sternum (59– 61). Stauros raised, narrow, tapering towards the valve face-mantle junction and widening 425 significantly on the valve mantle towards the mantle edge (Figs 59  $\&$  61). Central raphe endings straight, elongated, terminating onto weakly developed, elongated and flattened helictoglossae (Figs 427 59 & 61). A flatly ended cylindrical knob present at the central nodule (Figs 59 & 61). Areolae externally occluded by cribra, appearing sunken, especially close to the raphe-sternum (Figs 60 & 61). Terminal raphe endings terminating onto prominent helictoglossae within an expanded and thickened pore-free area corresponding to the curvature of the external silica flaps (Fig. 60). Several small areolae present at the end of the curved thickening (Fig. 60, arrowheads).

#### Page 95 of 113 and 12 and 13 and

 Cingulum composed of numerous open copulae bearing up to five rows of cribrate squarish or elongated areolae, ca. 38*–*45 in 10 μm (Figs 55, 59 & 62). Advalvar part of valvocopula pore-free beside the stauros (Fig. 59).

## **Taxonomic remarks**

 The morphological character pattern in *Craspedostauros macewanii* is most similar to *C. australis*  and *C*. *capensis* Cox. The three species share several features such as the presence of a bow tie- shaped fascia, rudimentary lip-like silica flaps extending from the raphe-sternum and partially covering the external central raphe endings, valve margin straight at the centre, and internally, a single knob at the central nodule (Table 1). Moreover, valve dimensions of *C. macewanii* (26–51 μm long, 4.5–5.5 μm wide) overlap with those reported for *C. australis* (35–78 μm long, 4–6 μm wide) and *C. capensis* (25–35 μm long, 4.5–5.5 μm wide). In *C. macewanii*, however, the stria density (28–31 in 10 μm) is significantly higher than in *C. capensis* (~19 in 10 μm) and lower than in *C. australis* (35 in 10 μm). In addition, *C. macewanii* can be distinguished from both *C. australis* and *C. capensis* by the presence of a distinct valve face-mantle junction running as a narrow, though clearly visible, pore-free ridge from apex to apex. *Craspedostauros macewanii* differs further from *C. capensis* in possessing areolae of a similar size throughout the entire valve (variable in *C. capensis*), and from *C. australis* in having convergent stria at the apices (parallel in *C. australis*) and extended apical hyaline zone (Cox 1999). The new taxon is also the only *Craspedostauros* species with girdle bands perforated by up to five rows of squarish areolae instead of two rows of usually transapically elongated areolae observed in other species.

 HOLOTYPE: Permanent slide SANDC-ST242 (prepared from sample ST242) deposited in the South African Diatom Collection housed by North-West University, Potchefstroom, South Africa.

456 TYPE LOCALITY: uShaka Sea World, Durban, South Africa (29° 52′ 02.79″ S, 31° 02′ 45.29″ E). Collected from the carapace of a captive juvenile loggerhead named "Bubbles" by R. Majewska, 28

June 2019.

ETYMOLOGY: The epithet honours Tony McEwan, the uShaka Sea World director, whose

 scientific enthusiasm and support to the sea turtle diatom project are highly appreciated and acknowledged.

 ECOLOGY: Epizoic on skin and carapaces of captive loggerheads and green turtles. Attaching to the animal surface through one end of the valve, motile in culture.

 The taxon was found on two captive loggerheads (a juvenile named "Bubbles" and an adult female named "DJ") and two captive green turtles (a subadult named "Calypso" and an adult male named "Napoleon") each time reaching relative abundance of 0.5–1%. All carapace samples containing *C. macewanii* were dominated by the so-called "marine gomphonemoids": *Poulinea* spp. and *Chelonicola* spp., accompanied by *Amphora* spp., *Nitzschia* spp., *Achnanthes elongata* and *A. squaliformis* Majewska et Van de Vijver, whereas the most abundant taxa in the four skin samples were *Tursiocola* spp., *Medlinella* sp., and the two previously mentioned *Achnanthes* species.

## *Craspedostauros alatus* **Majewska & Ashworth (Figs 63–74)**

 *Craspedostauros alatus* was found on the carapaces of several loggerhead sea turtles sampled at the Marine Turtle Rescue Centre in Pula, Croatia. The taxon co-occurred with *C. legouvelloanus*. As in the case of the latter, relative abundance of *C. alatus* was low (ca. 1–3% of the total diatom number). The observed morphological features of the Adriatic population agreed with the original description of the species (Majewska et al. 2018; Figs 63–74, Table 1). The examined specimens were 26*–*34 μm long and 3*–*5 μm wide (length/width ratio: 6.3*–*8.8), with stria density 24*–*27 in 10  $\mu$ m ( $n = 20$ ), and possessed all species-specific features, including a very distinct valve face-mantle

#### Page 97 of 113 and 12 and 13 and 13 and 13 and 13 and 14 and 15 and 16 and 16 Phycology

 junction and deep mantle (Figs 68, arrows, 69–71), wing-like silica flaps at the apices (Fig. 70), and 481 rectelevatum with central cavity (Figs 73 & 74).

### *DNA-based phylogeny*

 The genus *Craspedostauros* is monophyletic based on DNA sequence data generated from cultured material thus far (Fig. 75), though not with strong bootstrap support (bs < 50%). Regarding the taxa described here, *Craspedostauros macewanii* is sister to the rest of the clade (except *C. amphoroides*) with high support (bs = 96%), while *C. danayanus* is sister to *C. alyoubii* and *C. paradoxus* (bs =  $71\%$ ).

 Consistent with other molecular phylogenetic studies which include the genus (Ashworth et al. 2017), the position of the *Craspedostauros* clade can be found in a poorly supported (bs < 50%) assemblage containing the *Staurotropis* clade and a clade of marine *Achnanthes* species. This assemblage can be found within a clade with the Bacillariales (Supplementary Figure S1), though the relationship between the *Staurotropis*+*Achnanthes*+*Craspedostauros* clade and the three Bacillariales clades is poorly resolved. For taxa, strain voucher ID and GenBank accession numbers for strains used in the analysis see Supplementary Table S1.

## **DISCUSSION**

 The three new species described in the current study share most of the morphological characters typical of the genus *Craspedostauros*, such as squarish or rectangular areolae occluded by cribra on the valve and girdle bands, multiple copulae with at least two rows of perforations, and two fore and aft chloroplasts. Their linear or linear-lanceolate valve outline and the central constriction of the cell seen in girdle view resemble previously described species. Interestingly, two of the novel species, *C. macewanii* and *C. legouvelloanus*, present features not yet observed in any other member of the

 genus. The former possesses more than two rows of cribrate areolae on the girdle bands, whereas the latter shows shallow perforated septa. Moreover, the leatherback-associated *C. danayanus* presents a complete reduction of the stauros being the second, after *C. paradoxus*, *Craspedostauros* species lacking this character.

 It is interesting to note that as the number of character states, such as the reduction/loss of the stauros (*C. paradoxus* and *C. danayanus*) or addition of septate copulae (C. *legouvelloanus*), within *Craspedostauros* changes, the molecular data remain constant in their support (however tenuous) of monophyly for the genus. Cox (1999) ascribed the constricted girdle view to the presence of stauros. Yet the frustules of the two species lacking the latter, still show the central constriction, which may indicate that the lack of stauros is a secondary loss. One of the morphological features of the genus which has been maintained, regardless of newly described diversity, has been the cribrate areolar covering. While the degree of cribrum poration might change among species, the overall gestalt ultrastructure remains unchanged. Even more interesting is that this cribrum ultrastructure is also seen in *Staurotropis* and the *Achnanthes* species, which are commonly found (again, somewhat tenuously) sister to the *Craspedostauros* clade in molecular phylogenies. While there are other morphological similarities between the three genera, such as the stauros (though missing in some species of *Craspedostauros* and *Achnanthes*) and the fore and aft H-shaped or plate-like chloroplasts (missing in *Staurotropis* and some species of *Achnanthes*), so far it is the cribrate areolae ultrastructure that remains constant. In this context, the phylogenetic position of the genus *Druehlago*, which shares the same cribrum ultrastructure and the same chloroplast morphology of *Staurotropis* and *Achnanthes longipes* Agardh, but thus far lacks a stauros-bearing taxon, is all the more intriguing.

 Microscopical analyses of the fresh and critical-point-dried sea turtle skin pieces and barnacles revealed the mode of attachment and growth form of *C. danayanus* that attaches to the animal substratum through one pole of the cell. A similar mode of attachment to the natural substratum was

#### Page 99 of 113 and 12 and 12 and 13 and 14 and 15 and 16 Phycology

 observed in several members of the genus (R.Majewska, pers. observ.) suggesting that these taxa can either develop as firmly attached, sessile colonies or remain motile in less favourable conditions (e.g. in culture tubes).

 In the course of the on-going surveys on sea turtle-associated diatoms, a recently described taxon, *C. alatus*, was observed growing on the carapaces of several loggerhead sea turtles rescued in Croatia. *Craspedostauros alatus* was originally described from museum specimens of juvenile Kemp's ridleys (*Lepidochelys kempii* Garman) and a juvenile green turtle found cold-stunned and beyond recovery on the New York (USA) beaches during various seasons between 2012 and 2014 (Majewska et al. 2018). Although the relative abundance of *C. alatus* did not exceed 5.5% (current study, Majewska et al. 2018), observations of this taxon on a sea turtle from the Adriatic Sea may indicate that a) *C. alatus* is not an uncommon element of the sea turtle diatom flora; b) being associated with highly migratory animals such as sea turtles its geographical range is likely linked to that of its hosts.

 A similar conclusion can be drawn based on the records of *C. legouvelloanus*. The species occurred on several of the Adriatic loggerheads as well as on dozens of sea turtles belonging to the same species and their associated barnacles sampled on the eastern coast of South Africa. Even though the taxon was found in two different ocean basins, it cannot be excluded that the sea turtles acted as vectors that facilitated its dispersal among the various seas and oceans. There is a strong observational and molecular evidence that the Indian Ocean loggerheads interact and mate with the Atlantic members of the species (Bowen et al. 1994, Bowen and Karl 2007, Le Gouvello du Timat et al., in prep.). Thus, it is conceivable that any diatom able to endure the changing conditions during the migrations of their hosts and survive in competition with native flora would inoculate all appropriate and available media and substrata encountered. With the exception of *C. danayanus*, the sea turtle-associated *Craspedostauros* species, although common on the sea turtle carapaces, were never among the dominant taxa, and it is still unclear whether the animal body surface is their

#### Journal of Phycology **Page 100 of 113**

 preferred or alternative habitat. It is possible that the occurrence of these species in the sea turtle biofilm samples is linked to the presence of some other sea turtle epibionts (e.g. barnacles, sponges, bryozoans). *Craspedostauros danayanus* dominated most of the leatherback skin and barnacle samples that were analysed, and it is likely that this taxon is highly adapted to the conditions provided by the smooth body of the largest among the sea turtles, and, being associated with both the skin and the leatherback-specific barnacle species, *Platylepas coriacea*, its relationship with the host may be obligatory. Leatherbacks, contrary to other extant sea turtles, show the fully oceanic developmental pattern spending most of their lives in highly homogenous open-water environment devoid of refugia (Bolten 2003). They are unique among modern reptiles in being endothermal (Frair et al. 1972). This ability allows them to survive in both tropical and near-freezing waters (James et al. 2006). They are also significantly faster swimmers and deeper divers than other sea turtles (Eckert 2002, Doyle et al. 2008). Therefore, microhabitats provided by these animals, and thus their microbiomes, would differ substantially from those present on other sea turtles. Under such unique conditions, far from the diverse, species-rich shallow-water ecosystems, specific eco- physiological adaptations may be required to survive, and fewer diatom species would manage to thrive on the demanding substratum. An analogous phenomenon is known from marine cetaceans that seem to be colonised by only a few, highly specialized diatom taxa (e.g. Nemoto 1956, Holmes et al. 1993, Ferrario et al. 2018).

## **ACKNOWLEDGEMENTS**

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# Page 101 of 113 and 12 and 13 and 14 and 15 and 16 Phycology



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## **Figures legends**

 **Fig. 1**. Sampling locations where *Craspedostauros danayanus* (1), *C. legouvelloanus* (2), *C. macewanii* (3), and *C. alatus* (4) were found.

 **Figures 2–11.** *Craspedostauros danayanus*. **Fig. 2.** Living cells of *C. danayanus* and *Cylindrotheca* sp. attached to the leatherback skin scutes (light microscopy). **Fig. 3.** Stained colony of *C. danayanus* and associated bacteria on the leatherback skin scutes. **Fig. 4.** Valve view of a living cell (cultured strain). **Fig. 5.** Girdle view of a living cell (cultured strain). **Figs 6–11.** Scanning electron micrographs of *C. danayanus* attached to its original substratum. **Fig. 6.** Monospecific colony growing among the flaking skin of leatherback (dorsal side of the hind flipper). **Fig. 7.** Extremely delicate and fragile cells of *C. danayanus* attached to the leatherback skin (dorsal side of the hind flipper). **Fig. 8.** An overview of the leatherback-associated barnacle, *Platylepas coriacea*, colonized by *C. danayanus*. **Fig. 9.** A detail of the external part of the barnacle with a sheath of host sea turtle tissue overgrown with *C. danayanus*. Arrows indicate some of the monospecific clumps of *C. danayanus* colonies. **Fig. 10.** A detail of the moveable plates of the barnacle overgrown with *C. danayanus*. **Fig. 11.** A single cell of *C. danayanus* among dense colony of *Cylindrotheca* sp. attached to the folds in the moveable plates of *P. coriacea*. Scale bars: 10 µm = **Figs 3–5, 7, 11**; 50 µm = **Fig. 2**; 100 µm = **Figs 6, 9 & 10**; 1mm = **Fig. 8**

 **Figures 12–24.** *Craspedostauros danayanus*. **Figs 12–16.** Valve view (light micrographs). Arrows indicate the barely noticeable valve margins. **Figs 17–24.** Scanning electron micrographs. **Fig. 17.**  Detail of the apical part of the valve (external view). Arrowheads indicate the large irregular depression at the fold of the apical silica flap. **Fig. 18.** Frustule with partially detached girdle bands (external view). Arrowheads indicate the large irregular depression at the fold of the apical silica

## Page 109 of 113 and 12 Journal of Phycology

 flap. **Fig. 19.** Detail of the central part of the valve (external view). **Fig. 20.** Internal valve view. **Fig. 21.** Detail of the central part of the valve (internal view). **Fig. 22.** Cribrate areolae (internal view). **Fig. 23.** Detail of the apical part of the valve (internal view). Arrowheads indicate the asymmetrical thickening extending from the apical part of the raphe-sternum towards the valve margin. **Fig. 24.** Detail of the girdle bands. Scale bars: 10 µm = **Figs 12–16, 18, 20**; 1 µm = **Figs 17, 19, 21–24**

 **Figures 25–40.** *Craspedostauros legouvelloanus*. **Figs 25–30.** Light micrographs. **Figs 25, 26, 28– 30.** Girdle view. **Fig. 25.** Valve with two girdle bands attached. **Figs 28 & 29.** Frustules with detached valves. **Figs 26 & 30.** Complete frustules. Arrows indicate the biarcuate valve margin. **Fig. 27.** Valve view. **Figs 31–40.** Scanning electron micrographs. **Fig. 31.** External valve view. Arrows indicate depressions at the apical flap fold. **Fig. 32.** Detail of the apical part of the frustule (external view). **Fig. 33.** Valve with attached girdle bands (girdle view). **Fig. 34.** Detail of the girdle bands (internal view). Arrowheads indicate the internal thickening (septum). **Fig. 35.** Valve with partially detached girdle bands (internal view). **Fig. 36.** Internal valve view. Arrowheads indicate the slight expansion of the stauros on the side corresponding to the external lip-like silica flaps. **Fig. 37.** Detail of the apical part of the valve (internal view). **Fig. 38.** Detail of the central part of the valve (external view). **Figs 39 & 40.** Detail of the central part of the valve (internal view). Arrowheads indicate the hollows in the stauros-adjacent virgae. Scale bars: 10 µm = **Figs 25–31, 33, 35 & 36**; 1 µm = **Figs 32, 34 & 37–39**; 500nm = **Fig. 40** 

**Figures 41–47.** *Craspedostauros legouvelloanus*. **Fig. 41.** Living cells in culture (light

microscopy). Arrows indicate the H-shaped chloroplasts with one lobe pressed against each valve, a

feature characteristic of the genus. **Fig. 42.** External valve view (wild population). **Fig. 43.** External

valve view (cultured strain). **Fig. 44.** Internal valve view (wild population). **Fig. 45.** Internal valve

 view (cultured strain). **Fig. 46.** Detail of a girdle band showing internal thickening (septum) with perforations. **Fig. 47.** A single girdle band (external and internal view). Scale bars: 10 µm = **Figs 41–45**; 1 µm = **Figs 46 & 47**

 **Figures 48–62.** *Craspedostauros macewanii*. **Figs 48–54.** Light micrographs. **Figs 48–51.** Fresh (unpreserved) material. **Figs 48 & 51.** Living cells. **Fig. 48.** Girdle view. **Fig. 51.** Valve view. **Figs 49 & 50.** Damaged cells in girdle view with the cell content (including plastids) spilling beyond the cell wall. **Figs 49.** Arrow indicates the straight valve margin. **Figs 52–54.** Cleaned material. Detached valves in valve view. Arrows indicate the distinct valve face-mantle junction. **Figs 55–62.**  Scanning electron micrographs. **Fig. 55.** External valve view. **Fig. 56.** Detail of the apical part (external valve view). **Fig. 57.** Detail of the central area (external valve view). **Fig. 58.** Detail of the apical part (external girdle view). **Fig. 59.** Internal valve view and partially detached valvocopula. **Fig. 60.** Detail of the apical part (internal valve view). Arrowheads indicate several small areolae present at the end of the curved thickening. **Fig. 61.** Detail of the central area (internal valve view). **Fig. 62.** Detail of the valvocopula (internal view).

Scale bars: 10 µm = **Figs 48–55 & 59**; 1 µm = **Figs 56–58 & 60–62**

 **Figures 63–74.** *Craspedostauros alatus* (Adriatic population). **Figs 63–68.** Light micrographs. **Figs 63, 66 & 67.** Valve view. **Fig. 63.** Broken frustule with both valves lying in valve view. **Fig. 64.**  Single valve with attached girdle bands. **Figs 65 & 68.** Girdle view. Arrows indicate the clear valve face-mantle junction. **Figs 69–74.** Scanning electron micrographs. **Fig. 69.** Frustule with partially detached girdles bands (external view). **Fig. 70.** Detail of the apical part of the frustule with the winged-liked silica flaps, a feature typical of the species (external view). **Fig. 71.** Frustule with partially detached girdles bands (external girdle view). **Fig. 72.** Internal valve view. **Figs 73 & 74.**

 Detail of the central part of the valve (internal view). Scale bars: 10 µm = **Figs 63–69, 71 & 72**; 1 µm = **Figs 70 & 73**; 500 nm = **Fig. 74**

 **Figure 75.** Maximum likelihood (ML) phylogram based on the 3-gene dataset (nuclear-encoded ribosomal SSU, chloroplast encoded rbcL, psbC markers). For clarity, only the clade of raphid diatoms containing *Staurotropis*, *Craspedostauros*, and *Achnanthes* is presented in the figure. The ML tree presenting the complete taxon sampling can be viewed in the Supplementary Figure S1. 

## **Supplementary Figure S1**

 Maximum likelihood tree based on the 3-gene dataset (nuclear-encoded ribosomal SSU, chloroplast-encoded rbcL, psbC markers) with bootstrap values from 1000 pseudoreplicates over the corresponding nodes. The araphid pennate taxon outgroup *Asterionellopsis socialis* was used as the outgroup.



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Journal of Phycology **Page 114 of 113** 



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