### **ORIGINAL PAPER**



## *Cocconeis vaiamanuensis* sp. nov. (Bacillariophyceae) from Raivavae (South Pacific) and allied taxa: ultrastructural specificities and remarks about the polyphyletic genus *Cocconeis* Ehrenberg

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Received: 30 April 2020 / Revised: 21 October 2020 / Accepted: 10 December 2020 / Published online: 17 March 2021 © Senckenberg Gesellschaft für Naturforschung 2021

### Abstract

A small marine monoraphid diatom with a linear-elliptic shape and simple terminal raphe endings was present on Raivavae (South Pacific). Particularly due to its different stria structure on both valves, this taxon is here classified as *Cocconeis*. The new species was present on the rocky intertidal shore of the coral-reef lagoon, as an epiphyte on a turf. *Cocconeis vaiamanuensis* sp. nov. can be compared to some other monoraphids with a simple raphe system and a rod-like shape. The new taxon has small marginal processes on the sternum valve (SV) mantle, as previously reported for *Cocconeis peltoides*. Such processes were also previously observed in *Platessa* and *Psammothidium*. A cladistic analysis based on ultrastructure shows an affiliation between several close-by taxa. *Cocconeis* of the *C. peltoides* section are close to *Psammothidium*, whereas other *Cocconeis* without processes are closer to *Platessa* and *Achnanthidium*. A clone of *Cocconeis* cf. *sigillata* (SZCZCH1200) allowed for a molecular phylogeny to be reconstructed. The molecular signature of *Cocconeis* cf. *sigillata* is close to that of *Lemnicola hungarica*. *Cocconeis* is a genus with different and complex morphologies that may be split into independent clades (genera). The SV processes may be a vestigial character reminiscent of an ancestral state.

Keywords Ancestral character · Cladistic analysis · Marginal processes · Phylogenetic analysis · rbcL · Shape convergence

Communicated by B. Beszteri

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### Introduction

Raivavae (in Tahitian: Ra'ivāvae), part of the Austral Islands, also referred to as the Tupua'i Islands (South Pacific, Fig. 1), was first mentioned by Europeans in 1775, i.e., by the Spanish naval officer Tomas Gavagos, who was the commander onboard the frigate Aguila. Raivavae is a volcanic island (high island, erected between 10.6 and 5.4 Ma BP, Montaggioni 2015) with a large lagoon delineated by a coral-reef barrier with several motu (low coral islands), among which motu Vaiamanu (Fig. 1). Raivavae has a rocky shore and sheltered small bays with sandy beaches of coral sand. Ancestral marae are present on several parts of the island, witness to a rich culture and cultural past. The intertidal habitats of the lagoon were investigated in October 2018 to study monoraphid benthic diatoms. Several areas were sampled, in particular the fringe of Ruatara rock ("Rocher de l'Homme," name referring to a Tahitian legend) (Fig. 1). Other sediments and seaweeds were also sampled



Fig. 1 Raivavae location in the Austral Archipelago (South Pacific), with location of Ruatara rock and motu Vaiamanu

on the North coast of Raivavae. The monoraphid diatoms in Raivavae include several genera (Table 1), upon which Achnanthidium Kützing, 1844, Amphicocconeis De Stefano & D Marino, 2003, Astartiella Witkowski, Lange-Bertalot & Metzeltin, in Moser et al. 1998, Cocconeis Ehrenberg, 1837, Madinithidium Desrosiers, Witkowski & Riaux-Gobin, in Desrosiers et al. 2014, Planothidium Round & Bukhtiyarova, 1996, Pseudachnanthidium Riaux-Gobin, in Riaux-Gobin and Witkowski 2015 and Scalariella Riaux-Gobin, in Riaux-Gobin et al. 2012 (Table 1). Among these genera, Cocconeis and

 Table 1
 Assemblage of Achnanthales from Raivavae (South Pacific)

Achnanthidium sp.

Achnanthidium glyphos Riaux-Gobin, Witkowski & Compère

Amphicocconeis cf. discrepans (AWF Schmidt) Riaux-Gobin, Witkowski, Ector & Igersheim

Amphicocconeis cf. rodriguensis Riaux-Gobin & Al-Handal

Amphicocconeis spp. (several species)

Astartiella sp.

Cocconeis cf. borbonica Riaux-Gobin & Compère

C. carinata Riaux-Gobin, Ector & Witkowski

C. convexa MH Giffen

C. coralliensis Riaux-Gobin & Compère

- C. coronatoides Riaux-Gobin & Romero
- C. cupulifera Riaux-Gobin, OE Romero Compère & Al-Handal
- C. dapalistriata Riaux-Gobin, OE Romero, Compère & Al-Handal
- C. cf. dirupta W Gregory
- C. distans W Gregory
- C. geometrica Riaux-Gobin, OE Romero, Compère & Al-Handal
- C. guttata Hustedt & Aleem
- C. margaritata Riaux-Gobin & Al-Handal
- C. mascarenica Riaux-Gobin & Compère
- C. cf. mascarenica Riaux-Gobin & Compère
- C. cf. meisteri Riaux-Gobin, Compère, M Coste, Straub & Taxböck
- C. molesta Kützing
- C. cf. molesta Kützing
- C. paucistriata Riaux-Gobin, OE Romero, Compère & Al-Handal
- C. peltoides Hustedt
- C. cf. pseudograta Hustedt
- C. pseudomarginata W Gregory
- C. scutellum Ehrenberg
- C. sigillata Riaux-Gobin & Al-Handal
- C. vaiamanuensis sp. nov.
- C. sp. 1
- Madinithidium flexuistriatum (Riaux-Gobin, Compère & Witkowski) Witkowski, Riaux-Gobin & Desrosiers

Planothidium cf. mathurinense Riaux-Gobin & Al-Handal

Planothidium spp. (several species)

Planothidium rodriguense Riaux-Gobin & Compère

Pseudachnanthidium sp.

Scalariella sp.

Amphicocconeis had the most remarkable species diversity. Several new Amphicocconeis will be detailed elsewhere (C Riaux-Gobin pers. comm.). We here focus on a small linearelliptical monoraphid taxon described as new: Cocconeis vaiamanuensis sp. nov. The new taxon is detailed with LM (light microscope) and SEM (scanning electron microscope) and compared with allied taxa. Some characters of the taxon, such as its valve shape and presence of small processes on the SV mantle (see description below), prompted us to establish a cladistic analysis, based on ultrastructure, giving some light about a possible connection among several genera such as: Psammothidium Bukhtiyarova & Round, 1996, Platessa Lange-Bertalot in Krammer & Lange-Bertalot, 2004, Lemnicola Round & Basson, 1997, Pauliella Round & Basson, 1997, Achnanthidium, Karayevia Round & Bukhtiyarova ex Round, 1998, Rossithidium Round & Bukhtiyarova, 1996 and Cocconeis. It can be first noted that, except for Cocconeis, the previously cited genera are mostly associated with freshwater, while the new taxon is a priori strictly marine. The morphological limits between some of the latter genera are particularly blurred, with incertitude about the particular position of several of them (see below). The use of SEM is often decisive in delineating taxa, and therefore strengthening the results of morphological cladistic analyses that permit to sort them. In the same manner, concerning the molecular phylogeny, without direct observation of cells under a microscope, it can be challenging to predict the identity of some diatoms as belonging to a particular genus, solely on their position in the molecular phylogeny (Frankovich et al. 2018).

The genus Cocconeis is diverse in valve shape, ornamentation, and copula structure, allowing for several sub-groups to be identified, such as the Section Alveolatae De Stefano & OE Romero (De Stefano and Romero 2005) characterized by a bilayered complex SV, and the Section characterized by a marginal row of simple SV processes and complex RV striation (Riaux-Gobin et al. 2015), comprising Cocconeis peltoides Hustedt, 1939. The present study particularly seeks to provide insights into the phylogenetic and cladistic position of members of Cocconeis with or without processes. Such processes were named "poroids" in Tudesque et al. (2016) for some Platessa species, with the comment: "This type of 'poroid' requires further investigation because it could be reminiscent of an ancestral portula." These structures were named "rimoportula" in Romero & Riaux-Gobin (2014) for Cocconeis pseudograta Hustedt, 1939. Here we prefer to use the term "simple process", because these structures (that open externally via a tiny pore on the SV mantle and closed internally by a sort of polymorph domed hymen, see Riaux-Gobin et al. 2015) differ from true rimoportulae that are characterized by pair of lips in internal valve face, and simple pore or often more or less long tubes in external valve face, as illustrated in Round et al. (1990). Nevertheless, the difference in the complexity of these processes may be phenotypic differentiation among diatom Classes. SV processes were also

observed in several *Platessa* (Romero 2016; Wetzel et al. 2017), *Psammothidium* (Potapova 2012; Blanco et al. 2017), *Lemnicola* (Shi et al. 2018), *Rossithidium* (Potapova 2012), *Pauliella* (Round and Basson 1997), *Achnanthidium* (Jüttner et al. 2020), and *Karayevia* (Spaulding and Edlund 2008).

On the other hand, recent increases in the number of phylogenetic analyses have provided important clues allowing us to propose genetic affiliations. A *rbcL* molecular analysis of a clone of *Cocconeis* cf. *sigillata* SZCZCH1200 (collected in Laoshan Shangquan coastal area, China) (see *C. sigillata* Riaux-Gobin & Al-Handal, *in* Riaux-Gobin et al. 2011a), a taxon similar to *C. peltoides*, with marginal SV processes, allowed for the illustration of the complex and diverse relationships of *Cocconeis* (currently called *Cocconeis*) with other genera. Our results add to those of other recent phylogenetic investigations (i.e., Nakov 2014; Kulikovskiy et al. 2016, 2019; Thomas et al. 2016; Witkowski et al. 2016; Shi et al. 2018).

A polyphyletic position of the genus *Cocconeis* is considered, and the term Achnanthales (Order Achnanthales Silva, 1962), up to now treated as homogenous (De Stefano and Marino 2003; Le Cohu 2005) is here reconsidered, as previously addressed in several other studies (i.e., Cox 2006; Cox and Williams 2006; Kulikovskiy et al. 2016; Davidovich et al. 2017).

### Material and methods

### **Diatom materials**

Marine materials (sediments, seaweeds, diverse scrapings) collected in October 2018 from Raivavae (Austral Archipelago) were observed with LM and SEM, and are here illustrated and discussed. A strain of *Cocconeis* cf. *sigillata* was isolated and cultured by Chunlian Li at SZCZ (Szczecin University) from a sand beach in Laoshan Shangquan coastal area, China (36.092 N; 120.469 E), isolate SZCZCH1200; 101 stored in the Szczecin Diatom Culture Collection (SZCZ), University of Szczecin, collectors: Yu Shu-xian, Wang Yin-chu, Wang Xiu-jing, and Witkowski Andrzej, sampled in June 2015, Witkowski Lab Voucher SZCZCH1200. The molecular marker analysis was performed (see below), along with LM and SEM examination at SZCZ, Szczecin University.

### Sample preparation and examination

Materials (preserved in methanol) were washed with distilled water to remove salts (sedimentation method), treated with 30%  $H_2O_2$  for 2 h at 70 °C to remove organic matter, rinsed several times in distilled water, alcohol-desiccated, and mounted on glass slides using Naphrax. Diatom slides were examined with a Zeiss Axiophot 200, with phase contrast and differential interference contrast (Nomarski interference contrast) optics and photographed with a Canon PowerShot G6 digital camera

(CRIOBE-USR 3278, Perpignan, France). For SEM examination, drops of cleaned or raw material were filtered with a syringe-filter, through 1 µm Nuclepore® filters and rinsed twice with deionized (Milli-Q) water to remove salts. Filters were mounted onto aluminum stubs and air-dried before coating with gold-palladium alloy (EMSCOP SC 500 sputter coater) and examined with a Hitachi S–4500 SEM operated at 5 kV, calibrated with a Silicon grating TGX01 (C2M, Perpignan, France). Images kept in the authors' collection and handled with Photoshop and Adobe-illustrator. Measurements were performed using SEM images (60 individuals of *Cocconeis vaiamanuensis* sp. nov. were measured).

### **Cladistic analysis method**

A morphological cladistic analysis (ultrastructural characters) was applied to the taxa, including 16 characters (Table 2) and 34 taxa from 9 genera (Table 3). Taxa were selected following the accuracy of their SEM description, permitting or not to detail SV processes, RV transapical marginal poroids, and characteristics of the terminal raphe endings. The chosen characters consist of ultrastructural details allowing to define each of the nine genera included in the analysis: the shape of the valve, particular structures such as the mantle SV processes, valvocopula structure, stria pattern. For the outgroup, we selected a Planothidium (Planothidium juandenovense Riaux-Gobin & Witkowski in Riaux-Gobin et al. 2018). We used the software PAUP\*4.0a165 (Swofford 2003). We ran a Bootstrap method with a heuristic search algorithm to search for the most parsimonious trees (100 bootstraps). Characters were analyzed as unordered and unweighted. The resulting 100 most parsimonious trees were then used to build a 50% Majority-rule consensus tree, which is presented in Figure 8.

# Phylogenetic analysis method (DNA extraction, PCR, and molecular analysis)

Several clones, or strains, of *Cocconeis* and other genera, were cultured in SZCZ (Szczecin University), and molecular analyses were compared to data from GenBank®. Particularly, a clone of *Cocconeis* cf. *sigillata* SZCZCH1200 [see above details] was cultured and analyzed. Two strains of *Triparma pacifica* (Guillou & Chrétiennot-Dinet) Ichinomiya & Lopes dos Santos *in* Ichinomiya et al. (2016), p. 1432; Basionym: *Bolidomonas pacifica* Guillou & Chrétiennot-Dinet *in* Guillou et al. 1999, p. 371) were chosen as an outgroup. The final *rbcL* gene trees are available in Fig. 8. The full list of sequences used in phylogenetic analyses along with GenBank® accession numbers is available (Table 4).

DNA extraction using Chelex® 100 resin (Bio-Rad, cat. no. 142–2842-MSDS) followed the method described *in* Kryk et al. (2020). The molecular marker *rbc*L was amplified using primer and PCR protocol as described in Dąbek et al. (2017). The

sequencing of the PCR products was performed using BigDve Terminator v.3.1 chemistry and ABI3730 xl sequencer by the oligo.pl DNA Sequencing Laboratory at the Institute of Biochemistry and Biophysics, Polish Academy of Sciences, Warsaw, Poland. For sequence assembling, BioEdit ver. 7.2.5 (Hall 1999) was used. For maximum likelihood (ML) estimation, phylogenetic tree was inferred with 1000 bootstrap replicates using rapid bootstrap analysis in RAxML v.8.1 (Stamatakis 2014). The best score for ML tree was chosen as the final tree, and bootstrap support values (bv) were added at nodes. For Bayesian Interference, the tree was carried out using MrBayes v3.2.7 (Huelsenbeck and Ronquist 2001). The data was partitioned as three different codon positions. Two Bayesian inference analyses each with four chains (one cold and three heated) were run using GTR + G + I model. During this procedure, the standard deviation of split frequencies was around 0.01 when it reached 150,000,000 generations, which were run per analysis with sampling every 1000th iteration, generating in total of 150, 000 samples. 25% of the samples were discarded, then the rest were used to get a majority rule consensus tree and obtain posterior probabilities for nodes.

### Results

# Marine Achnanthales assemblage from Raivavae (Table 1)

The microphytobenthos from the intertidal environments of Raivavae is highly diversified (in term of species number), particularly within the monoraphids (Table 1). We note the presence of several pantropical taxa, from which several were previously reported from the Indian Ocean. Some taxa belonging to Olifantiella Riaux-Gobin & Compère, 2009, and Amicula Witkowski, Lange-Bertalot & Metzeltin, 2000, were also present. Most monoraphids (Table 1) are present in several other places in the South Pacific. In contrast, Cocconeis vaiamanuensis sp. nov. may be a local endemic, since it has never been cited from elsewhere. During several sampling campaigns in the South Pacific, we noticed different productivity and species diversity of benthic diatoms, particularly concerning the monoraphids, as a function of the geomorphology of the site: atolls (carbonate environments) versus high volcanic islands (siliceous environments, see remarks in Riaux-Gobin et al. 2014). More than 37 monoraphid taxa were reported from Raivavae, upon which 25 Cocconeis (Table 1). Amphicocconeis, with several taxa, is detailed elsewhere.

### **Cocconeis vaiamanuensis Riaux-Gobin, Witkowski** & Ector sp. nov. (Table 2; LM Fig. 2 a-p; SEM Figs. 3, 4, 5, 6)

Class: Bacillariophyceae Haeckel, 1878 Order: Achnanthales Silva, 1962 Family: Cocconeidaceae Kützing, 1844 Genus: Cocconeis Ehrenberg, 1837

### Description

Frustule solitary, valve oblong-elliptic to linear in the larger specimens (SEM, n = 60, length 7–15 µm, width 3–5 µm, L/W 2.2), with round apices. Never observed in girdle view during this survey.

Sternum valve (SV) slightly concave (Fig. 3), with a narrow, flat and straight sternum. Small round areolae (ca. eight per stria; 4-5 in 1 µm). SV striae uniseriate ( $n = 32, 16-22, \bar{x} = 19.6 \pm 1.3$  in 10 µm), parallel in mid-valve to radiate and slightly denser on apices. Striae composed of tiny areolae, more or less in zig-zag, internally closed by strongly convex hymenes without obvious slits or punctuations (Fig. 4b, c). Running all over the valve, a marginal embossed apical and hyaline area similar to a crista marginalis (Fig. 3, arrow). Mantle narrow with one row of small pores providing access to a process (Fig. 3f, arrows). This small process is internally closed by a hemispheric plug or hymen, slightly different and smaller in size than that of the SV areolae (Fig. 4d, arrowhead). One process regularly faces each stria (Fig. 4b, arrowheads). Strong virgae, externally embossed on their most marginal part (Fig. 3d, arrowheads). Relatively high cingulum composed of several open and large and thin cingulae devoid of ornamentation (Fig. 3, arrowhead, c, e). SVVC apparently with no fimbriae (Fig. 4c, arrowhead and ellipse). SV valve possibly bi-layered (Fig. 4d, framed arrowheads).

Raphe valve (RV) strongly convex (Fig. 5c). Striae radiate, slightly denser on apices  $(n = 30, 20-24, \bar{x} 21.3 \pm 1.2 \text{ in})$ 10 µm), composed of small areolae, biseriate near the axial area, uniseriate on a short median section, and up to quadriseriate near the margin. On rare specimens (one observed, see supplementary Fig. 11), the RV striae are distinctly biseriate, with alternate areolae. RV areola hymenes internally convex with a crenulated border giving it a star-like shape (Fig. 6d, e, arrowheads). A marginal row of dense and apically elongate-oblong poroids (60–72,  $\bar{x}$  65.2 ± 3.5 in 10  $\mu$ m, Fig. 5, black arrowhead), separated from the rest of the valve by a large hyaline apical area (Fig. 5, white arrowhead), corresponding to an internal flat unraised rim (Fig. 6). Central area reduced to absent (no evident fascia, Fig. 5, or short hemi-fascia on some specimens). Raphe filiform, straight. Proximal raphe endings externally close to each other (slightly bent on the same side, Fig. 5e) and internally bent in opposite directions (Fig. 6b). Terminal raphe endings simple, close to the margin (Fig. 5b, arrowhead), apically surrounded by an anchor-like silica fold (Fig. 5d, arrowhead) with no connection to the interior of the valve. No RV areolae on apices (Fig. 5b, ellipse). RVVC taking place on the border of the RV, with no fimbria (Fig. 6, arrowhead), or a slightly undulated edge (Fig. 6f, framed

 Table 2
 Characters, state, and coding used in the Cladistic Analysis. \*See definition in the text

N°	Definition	States	Coding
1	Frustule bent along apical axis	Not bent	0
		Bent	1
		Not detailed	2
2	Raphe terminal endings	Simple	0
		Simple with groove	1
		Short fissure slightly deflected	2
		Large fissure	3
3	Valve shape	Elliptic	0
		Subrostrate	1
		Round-elliptic	2
		Linear	3
		Oblong-elliptic	4
4	SV curvature	Flat	0
		Concave	1
		Convex or flat with a concave sternum	2
		Convex	3
5	RV curvature	Flat	0
		Concave	1
		Convex	2
6	Fascia	No Fascia	0
		Fascia on SV	1
		Fascia on RV	2
		Fascia on both valves	3
7	SV crista marginalis	Absent	0
	2	Present	1
8 9	Stria spacing	Regularly spaced	0
	Sum sprenng	Slightly denser on apices	1
		Markedly denser on apices	2
	RV stria orientation	Parallel to almost parallel	0
		Radiate	1
		Only strongly radiate on anices	2
	SV & RV structure	Similar	0
10	SV & RV subture	Closely similar	1
		Dissimilar	2
11	SV striae	Uniseriate	0
11	5 v Sulle	Uni to hiseriate	1
		Regularly hiseriate (or triseriate)	2
		Marginally multiseriate	3
		Multiseriate (>3)	4
12	RV striae	Uniseriate	0
12	it v suide	Uni to hiseriate	1
		Regularly hiseriate (or triseriate)	2
		Multiseriate (>3)	2
13	SV & BV strip density	Similar	0
15	5 V & ICV stila delisity	Closely similar	1
		Dissimilar	1
14	SV one row of processes on mantle*	Apparently abcent	0
14	SV one row or processes on manue	Present tiny, one per strip	1
		Denser than stripe	1
		Slit like one per strip	2
		Slit-like, one per sura	3
15	PV one row of obland paraids on mantle*	Apparently absent	7
13	Ky one low of obioing potoids on manue*	Apparenti y and nor stric	1
		Present tiny, one per suita	1
		Clit like one not this	2
		Slit-like, one per stria	3
16	V-loslas	Siit-like, denser than striae	4
10	vaivocopuiae	Apparentiy with straight to simply undulated edge	0
		r inoriae more or less complex	1
		not detailed	2



**Fig. 2** *Cocconeis vaiamanuensis* sp. nov. LM (Raivavae, Ruatara rock). **a–j** SV with linear-elliptical shape, round apices and regularly spaced striae; **k** RV with almost invisible striae; **l–p** complete frustules at different foci allowing for the RV and SV of each specimen to be

delineated and visualized. Scale bars = 10  $\mu$ m (Nomarski interference contrast); **q** *Cocconeis finmarchica* Grunow in Cleve & Grunow. Original drawings. Note the marginal hyaline area on both valves, and the RV fascia. Scale bar = 10  $\mu$ m.

Fig. 3 Cocconeis vaiamanuensis sp. nov. SEM (Raivavae, Ruatara rock). a-b SV in external view. Slightly concave, with regularly spaced striae. Note the marginal crista marginalis (arrow). Cingulum composed of several large copulae (arrowhead); c dense marginal striation of the RV (arrowhead); d marginal raised portion of the SV virgae (multiple arrowheads); e-f position of the small aperture of each process on the SV mantle (twin arrowheads, twin arrows). Scale bars =  $3 \mu m$ (c), 2 µm (a–b, d), 700 nm (e), 300 nm (f)



arrowheads). Central area almost absent (Fig. 5). Helictoglossa straight and low (Fig. 6g).

### Holotype:

Whole slide from the sample RAI 20, deposited at NHM (BM 81917 material). Holotype illustrated in Fig. 5.

### Type locality:

Ruatara rock (Rocher de l'Homme), Raivavae (Austral Islands, South Pacific), RAI 20 (intertidal red macroalgal turf on small fissures of the rock). Geo localization: S 23° 51.274'; W 147° 39.595'; T°C 23°.7, salinity < 40‰. Sampled by C. Riaux-Gobin on 08 10 2018.

### Ecology:

Relatively rare, marine intertidal taxon, living on rocky shore covered by turf. The new taxon was present in RAI 20 (see above) but also, in lower densities, on RAI 18 (same GPS position-geo localization, on red-dark short algae), and RAI 21 (same geo-localization, red-brown algae, longer than those in RAI 18). Taxonomy of the varied short macroalgae (turf) not available.

### **Etymology:**

The epithet *vaiamanuensis* refers to the most famous motu from Raivavae, the motu Vaiamanu (in Tahitian "the place where there are birds"), also named "motu Piscine."

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Fig. 4 Cocconeis vaiamanuensis sp. nov. SEM (Raivavae, Ruatara rock). a SV in internal view. Regular arrangement of striae composed of small round areolae with convex hymenes. Areolae near the sternum less dense than on the margin (arrowhead); a narrow and straight SV sternum; **b** SV processes, facing each stria, smaller than the areolae (multiple arrowheads); c broken SV with a large copula (arrowhead), SVVC still attached to the SV, with no fimbriae (ellipse); d Detail of the SV processes (arrowhead), note that the SV is possibly bi-layered (framed arrowheads). Scale bars = 2 µm (a), 1 µm (b-c), 300 nm (d)

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### Distribution:

Until now only found on Raivavae, as a possible local endemic [i.e., to the best of our knowledge, absent in Rapa (Austral Islands), as well as in the other visited Polynesian Archipelagos. Also absent from Mascarenes and Scattered Islands (Indian Ocean), and other tropical locations referenced, i.e., *in* Riaux-Gobin et al. (2011c, p. 7).

### **Remarks:**

Relatively rare taxon (< 5% of the all benthic diatom assemblage in RAI 20; occasionally present only on three samples in the vicinity of Ruatara). Some dissimilarity concerning the ultrastructure of the valves with *Cocconeis nugalas* MH Hohn & Hellerman (Hohn and Hellerman 1966) [synonym *C. hauniensis* Witkowski (Witkowski 1993; Desianti et al. 2015; see also SEM *in* Riaux-Gobin and Romero 2003, pl. 41–43]. The SV sternum is elliptical and slightly concave in *C. nugalas* while narrow-linear and flat in the new taxon. *Cocconeis nugalas* has no *crista marginalis*. The SV virgae are more significantly embossed in the new taxon than in *C. nugalas*. Furthermore, on *C. vaiamanensis* the RV striae are biseriate on a part of their length and composed of tiny areolae, while bigger and uniseriate on *C. nugalas*  (Riaux-Gobin and Romero 2003: pl. 43, Fig. 3). The RV of *C. vaiamanuensis* has a margin with almost four oblong poroids between each stria, seemingly more complex than in *C. nugalas* where there is only one marginal oblong poroid between each stria. The hyaline marginal area is narrower in *C. nugalas*. Furthermore, the valve shape is different between the two taxa: elliptical in *C. nugalas* and linear-elliptical to rod-like in the new taxon. The two taxa are thus significantly dissimilar.

Based on LM, similarities also exist with *Cocconeis* finmarchica Grunow in Cleve & Grunow (Cleve and Grunow 1880, p. 16, pl. 1, Fig. 1; illustrated here in Fig. 2q), also a small taxon, with a narrow and straight SV sternum and a SV hyaline marginal area (possibly a *crista marginalis*), and an RV hyaline margin. The RV of the latter has a slightly denser RV striation than the SV (24 in 10  $\mu$ m, 20 in 10  $\mu$ m in the SV) and radiate RV striae, such as in the new taxon. The only points that do not match the new taxon are the narrow and long fascia on the RV in *C. finmarchica*, and the shape of the frustule that is ellipsoid. No recent bibliography offers SEM for *C. finmarchica*.

For *Cocconeis* with SV processes, and concerning the taxonomic key presented in Riaux-Gobin et al. (2015), the addition of a supplementary group would be necessary to identify the new taxon: i.e., "Group 5 with SV striae composed of numerous areolae, with no partition, and with a narrow Fig. 5 Cocconeis vaiamanuensis sp. nov. SEM (Raivavae, Ruatara rock). a RV in external view. Whole valve with radiate striae. marginal hyaline area (white arrowhead), one row of marginal elongate poroids, denser than the striae (black arrowhead); b simple terminal raphe ending (arrowhead), absence of elongate poroids on apex (ellipse); c high cingulum (arrowhead); d anchorlike and raised fold around the simple terminal raphe ending (arrowhead); e proximal raphe endings small and bent on primary side (twin arrowheads); f RV striae with small areolae in quincunx ending in a pyramidal shape before the marginal hyaline area; g central area absentmissing. Scale bars =  $2 \mu m (\mathbf{a}, \mathbf{c})$ , 1 µm (b, e-f), 700 nm (g), 600 nm (**d**)



sternum void of areolae." This key would clearly benefit from references to the structure of the RV when available. However, neither *C. inaequalistriata* Riaux-Gobin, OE Romero, Compère & Al-Handal (Riaux-Gobin et al. 2011c, p. 28–29, pl. 48, Figs. 1–6; Riaux-Gobin et al. 2015, Fig. 13), nor *C.* sp4 *in* Riaux-Gobin et al. (2015), Figs. 12, 30–32; also illustrated *in* Riaux-Gobin et al. 2011c,? *Cocconeis* sp. 1, p. 41, pl. 85, Figs. 1–4) match the new taxon.

The genera *Psammothidium* and *Platessa* also have simple terminal raphe endings, but none correspond to our marine taxon. A cladistic analysis, based on ultrastructural characters (see below and Fig. 7), allows us better to understand the degree of proximity of genera and taxa. A phylogenetic tree including a taxon pertaining to the *Cocconeis peltoides* group also allows for some remarks to be made (see below and Fig. 8).

# Cladistic analysis based on ultrastructure (Tables 3, 4, Fig. 7)

It can be first pointed that several selected characters (Table 3) may appear as blurry, such as the valve shape (difficult to define accurately) or curvature of the frustule in cingular view (along the apical axis), a major-determinant character often neglected in descriptions. The morphology of the valvocopulae (with or without fimbriae), also an important character, is not often detailed.

*Planothidium juandenovense* is the outgroup, with groups of areolae with a multiseriate arrangement on the SV mantle and terminal raphe fissures strongly bent.

*Platessa bahlsii* Potapova (Potapova 2012) is the only taxon in the analysis showing multiseriate striae, seemingly belonging to *Planothidium* but with simple terminal raphe endings.

Fig. 6 Cocconeis vaiamanuensis sp. nov. SEM (Raivavae, Ruatara rock). a RV in internal view. Whole valve with a small portion of the RVVC, apparently with no fimbriae (arrowhead), low helictoglossa; b proximal raphe endings curved in opposite sides; c unraised marginal hyaline area (arrowhead); d domed RV areolae, in quincunx (arrowhead); e crenulated periphery of the RV hymenes (arrowhead); f probable edge of the RVVC slightly undulated, with no fimbriae (framed arrowheads); g detail of apex without poroids (ellipse). Scale bars =  $2 \mu m (\mathbf{a}, \mathbf{c})$ ,  $1 \mu m (\mathbf{b}, \mathbf{c})$ g), 600 nm (f), 200 nm (d-e)



*Platessa bahlsii* appears as an independent taxon (monophyletic). It would be recommended to culture such a taxon and to do molecular analyses to evaluate its degree of affiliation (proximity) to *Planothidium*. Kulikovskiy et al. (2020) recently suggested that "*P. bahlsii* belongs to the group of species without sinus or cavum in the genus *Planothidium*", and proposed the new combination *Planothidium bahlsii* (Potapova) Kulikovskiy, Glushchenko & Kociolek, *in* Kulikovskiy et al. (2020).

The cladistic analysis (Fig. 7) roughly has two poles: (1) Group I composed of taxa pertaining to *Platessa*, *Cocconeis* without SV processes and *Achnanthidium* and (2) Group II composed of *Psammothidium*, *Cocconeis* with SV processes and other atypic taxa upon which *Rossithidium* that is not accepted as different from *Psammothidium* by various authors (see Kulikovskiy et al. 2016; Jüttner et al. 2020).

Illustrating the difficulty of establishing limits between *Achnanthes* Bory, 1822, *Psammothidium*, *Platessa* and *Cocconeis*, we can cite the intricate taxonomic history of *Cocconeis therezienii* Le Cohu & R.Maillard ex Van de Vijver & Le Cohu (Van de Vijver and Le Cohu 2019). The interpretative analysis of a taxon by different authors may illustrate the difficulty of defining it correctly. For example, *Platessa oblongella* (Østrup) CE Wetzel, Lange-Bertalot & Ector (Wetzel et al. 2017) and its synonym *Karayevia oblongella* (Østrup) Aboal, *in* Aboal et al. (2003), do not appear at the same level in the cladistic analysis (Fig. 7), due to SEM observations that do not exactly match.

![](_page_11_Figure_2.jpeg)

![](_page_11_Figure_3.jpeg)

Fig. 7 Cladistic Analysis based on ultrastructure. OG = out group. See comments in the text

![](_page_12_Figure_2.jpeg)

Fig. 8 Phylogenetic tree, showing the position of different groups of Cocconeis. See comments in the text

n°	Taxon name, authorities	Character number			maracter										Authors - year, SEM documents			
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	
1	Achnanthidium atomoides O Monnier, Lange-Bertalot & Ector	1	0	4	3	1	2	0	2	2	2	0	0	2	1	1	2	Monnier et al. 2004
2	Achnanthidium minutissimum (Kiitzing) Czermeglei	1	2	3	3	1	2	0	2	2	0	0	0	1	3	1	0	Czarnecki 1994, Potapova 2009
3	Achanthidium parallelum JR Carter ex Jüttner, DM Williams, Ector & CF Wetzel	1	0	3	2	1	2	0	0	0	1	0	0	1	3	3	0	Jüttner et al. 2020
4	Achanthidium petersenii (Hustedt) CE Wetzel, Ector, DM Williams & Jüttner (Rossithidium)	2	0	4	2	1	2	0	0	1	1	0	0	1	3	3	0	Jüttner et al. 2020
5	Achnanthidium subhudsonis var. kraeuselii (Cholnoky) Cantonati & Lange-Bertalot	1	3	4	1	2	0	0	2	1	2	0	0	2	1	1	2	Kusber et al. 2017, Spaulding and Potapova 2014
6	Cocconeis alucitae Riaux-Gobin & Compère	0	0	2	3	0	0	0	0	1	2	3	1	1	0	0	1	Riaux-Gobin and Compère 2008
7	Cocconeis coronatoides Riguy-Gobin & Romero	0	0	0	3	1	0	1	0	1	2	1	0	2	0	0	1	Riaux-Gobin et al. 2010, 2011b
8	Cocconeis germainii Riaux-Gobin, Witkowski & OE Romero	0	0	0	2	0	0	0	0	1	2	0	0	2	1	4	0	Riaux-Gobin et al. 2007
9	Cocconeis neuquina Frenguelli	0	0	2	3	0	0	0	0	1	2	0	0	2	0	0	1	Frenguelli 1942, García et al. 2018
10	Cocconeis nugalas MH Hohn & Hellerman	0	0	2	2	0	0	0	0	1	2	0	0	0	1	4	0	Hohn and Hellermann 1966, Riaux-Gobin et al. 2015
11	Cocconeis peltoides Hustedt	0	0	0	2	1	0	1	0	1	2	0	0	2	1	4	0	Hustedt 1939, Riaux-Gobin et al. 2011a
12	Cocconeis pseudograta Hustedt	0	1	2	1	2	0	1	0	1	2	0	0	2	1	4	0	Hustedt 1939, Romero and Riaux-Gobin 2014
13	Cocconeis sigillata Riaux-Gobin & Al-Handal	0	0	0	2	1	0	1	0	1	2	2	0	2	1	4	0	Riaux-Gobin et al. 2011a
14	Cocconeis vaiamanuensis Riaux-Gobin, Witkowski & Ector	0	1	4	1	2	0	1	0	1	2	0	1	0	1	4	0	present paper
15	Karayevia oblongella (Østrup) Aboal	0	0	4	0	2	2	0	0	1	2	0	0	2	1	1	2	Aboal et al. 2003
16	Lemnicola hungarica (Grunow) Round & Basson	0	3	1	3	1	2	0	0	0	2	2	2	0	1	1	0	Round and Basson 1997, Shi et al. 2018
17	Lemnicola uniseriata Y Shi & B-H Kim	0	3	0	3	3	3	0	0	1	0	0	0	0	1	1	2	Shi et al. 2018
18	Pauliella taeniata (Grunow) Round & Basson	2	0	3	0	0	0	0	0	0	0	0	0	0	3	3	2	Round and Basson 1997
19	Planothidium juandenovense Riaux-Gobin & Witkowski (OG)	0	3	0	2	2	0	1	0	1	0	4	3	0	0	0	2	Riaux-Gobin et al. 2018
20	Platessa bahlsii Potapova	0	0	0	2	2	0	0	0	1	0	4	3	0	0	0	2	Potapova 2012
21	Platessa brevicostata (Hustedt) Lange-Bertalot	0	0	4	0	1	0	0	0	1	1	2	2	0	1	1	0	Krammer and Lange-Bertalot 2004, Romero 2016
22	Platessa cataractarum (Hustedt) Lange-Bertalot	0	0	0	1	0	2	0	0	1	2	1	1	0	1	1	0	Krammer and Lange-Bertalot 2004, Romero 2016
23	Platessa guianensis Le Cohu, Tudesque & CE Wetzel	1	0	0	3	1	2	0	0	1	2	2	2	1	3	1	0	Tudesque et al. 2016
24	Platessa itoupensis Tudesque, Le Cohu & CE Wetzel	0	1	0	3	1	0	0	1	1	1	2	2	1	0	0	0	Tudesque et al. 2016
25	Platessa kingstonii Potapova	0	0	4	0	0	2	0	0	1	2	2	0	2	0	1	0	Potapova 2012
26	Platessa oblongella (Østrup)	1	0	0	3	1	2	0	0	1	2	1	0	2	1	2	0	Wetzel et al. 2017, Potapova 2011
	CE Wetzel, Lange-Bertalot																	· •
	& Ector																	

 Table 3
 Taxa (34 species), character coding (16 ultrastructural characters), author and year, where can be found the SEM documents referring to each taxon). OG = out group

### Table 3 (continued)

n° Taxon name, authorities			Character , , , , , , , , , , , , , , , , , , ,														Authors - year, SEM documents	
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	
27	<i>Platessa saxonica</i> (Krasske ex Hustedt) CE Wetzel, Lange-Bertalot & Ector	1	0	4	2	1	2	0	0	1	2	2	0	2	1	1	0	Wetzel et al. 2017
28	Platessa stewartii (RM Patrick) Potapova	0	0	0	2	0	3	0	0	1	0	2	2	0	1	1	0	Potapova 2012
29	Psammothidium curtissimum (JR Carter) Aboal	0	0	0	0	0	3	0	0	1	0	0	0	0	2	2	0	Aboal et al. 2003, Potapova 2010, Blanco 2016
30	Psammothidium harveyi (Reimer) Potapova	0	2	1	1	2	3	0	0	1	0	0	0	0	1	1	0	Potapova 2012
31	Psammothidium levanderi (Hustedt) Bukhtiyarova & Round	0	0	0	1	0	0	0	0	1	2	0	0	0	3	1	2	Bukhtiyarova and Round 1996, Blanco et al. 2017
32	Psammothidium semiapertum (Hustedt) Aboal	0	0	0	2	2	3	0	0	1	0	0	0	0	2	2	2	Aboal et al. 2003, Potapova 2012
33	Psammothidium toroi S Blanco, Pla-Rabes, CE Wetzel & I Granados	0	0	0	1	2	0	0	0	1	2	0	0	0	2	2	2	Blanco et al. 2017
34	Rossithidium anastasiae (Kaczmarska) Potapova	0	0	3	0	0	0	0	0	0	0	0	0	0	1	1	2	Potapova 2012

### Molecular phylogenetic position of *Cocconeis cf.* sigillata SZCZCH1200 and some other *Cocconeis* (*Figs.* 8, 9)

Our rbcL-based phylogenetic trees recover monoraphid diatoms as polyphyletic and spreads them over two diatoms clades: *Achnanthes* + Bacillariaceae sister to Cocconeidaceae + other monoraphid and raphid diatoms (100% ML bootstrap, 0.5536 BI posterior probability), although the support of some branches within the two clades is very low.

As mentioned, *Achnanthes* as representative of Achnanthaceae is placed within the large clade together within Bacillariaceae family as sister to *Nitzschia* Hassall, 1845, *Psammodictyon* DG Mann, *in* Round et al. 1990 and *Tryblionella* W Smith, 1853, however, with a low support (< 50% ML bootstrap, 0.5035 BI posterior probability). This position of *Achnanthes* spp. is stable on either single gene (Mann et al. 2020), three genes (Theriot et al. 2015; Ashworth et al. 2017) or in multigene (Theriot et al. 2010; Nakov et al. 2018a, b) trees.

Cocconeidaceae, represented in our phylogeny by several strains of *Cocconeis* Ehrenberg, are spread over a few groups and are seemingly polyphyletic. The major *Cocconeis*-bearing clade is recovered fairly isolated and is positioned at the base of the clade containing all monoraphids and raphids, excluding Bacillariaceae. Here, *Cocconeis pediculus* Ehrenberg, 1838, and *C. placentula* Ehrenberg, 1838, create a fairly distant and monophyletic group, which is in a sister relationship to *Cocconeis* sp. KC309551 (strain name MPA2013ECT3901, see Table 4) (81% ML bootstrap, 1 BI posterior probability),

and together they are grading into a smaller group composed of *Berkeleya hyalina* (Round & ME Brooks) EJ Cox, 1975 (< 50% ML bootstrap) and two *Planothidium* and *Cocconeis* cf. *cupulifera* Riaux-Gobin, OE Romero, Compère & Al-Handal, 2011c, KT943680 (strain name SZCZCH662) (< 50% ML bootstrap). The relationship of the latter taxon to *Planothidium* spp. is moderate (65% ML bootstrap, 1 BI posterior probability).

Psammothidium, Rossithidium and some Achnanthidium form a paraphyletic group with low support at the nodes but usually moderate to high (>60% ML bootstrap, >0.68 BI posterior probability) in the particular clades. This clade is grading further into a monophyletic branch composed of Achnanthidium (10 species), and another clade of Pauliella taeniata (Grunow) Round & Basson, 1997, both with moderate to high support. Cocconeis cf. sigillata MT015687 (strain name SZCZCH1200) and C. cf. mascarenica Riaux-Gobin & Compère, 2008 KT943679 (strain name SZCZCH283) (73% ML bootstrap, 0.9999 BI posterior probability) are nested within this clade (though, with low by), with unresolved trichotomy including Lemnicola hungarica (Grunow) Round & Basson, 1997, and Cocconeis sp. KT943614 (strain name SZCZP67) (here with also very low bv). Sister to the above monoraphid clades is *Planothidium* related to the clade of Cymbellales, however, with a low support (54% ML bootstrap, 0.9812 BI posterior probability).

*Cocconeis stauroneiformis* Okuno, 1957, AB430694 (strain name s0230) is positioned within Surirellales and hence fairly distant from the above monoraphids

### Table 4All strains used for the phylogenetic analysis, including the taxon name, strain ID, and assession number (*rbc* L) in genbank

Taxon	Strain ID	Assession number ( <i>rbc</i> L)	Taxon	Strain ID	Assession number ( <i>rbc</i> L)	
Achnanthes coarctata	UTEX FD185	HQ912458	Fragilariopsis cylindrus	E8C2	EF423499	
Achnanthes sp.	SZCZCH113	KT943615	Fragilariopsis kerguelensis	E13B2	EF423500	
Achnanthes sp.	MPA-2013 SanNic1Achnan	KC309545	Geissleria decussis	FD50	KJ011830	
Achnanthes sp.	MPA-2013 ECT3911Achnan	KC309547	Hantzschia amphioxys var. major	A4	HQ912390	
Achnanthes sp.	SZCZM119	KT943616	Karayevia ploenensis var. gessneri	D03_034	KM084931	
Achnanthes sp.	MPA-2013 ECT3883Achnan	KC309546	Kolbesia sinica	SZCZM123	KT943677	
Achnanthidium catenatum	TCC849	KY799133	Lemnicola hungarica	UTEX FD456	HQ912490	
Achnanthidium cf. lineare	NA-2016 strain B397	KR709273	Lemnicola hungarica	Lemn1	KJ658388	
Achnanthidium coarctatum	UTEXFD185	HO912458	Madinithidium vietnamica	SVN252	MH231749	
Achnanthidium daonense	PS3	KJ658395	Mastogloia aquilegiae	KSA2015-49 Masto-B1	MH064100	
Achnanthidium digitatum	SPITS EBA-A 22	KU687461	Mastogloia sp	29×07-6B	HO912496	
Achnanthidium digitatum	SPITS M2AnlusB 32	KU687471	Mastogloia sp	SA17 Masto-C1	MH064101	
Achnanthidium kranzii	A100r	K 1658379	Mastogloia sp	KSA2016-44 Masto-A37	MH064102	
Achnanthidium	TCC746	KF050640	Navicula cryntocenhala	LITEX ED100	HO012467	
minutissimum	TCC740	KI 7570124		TA 105	NV220200	
minutissimum	100/48	K¥/99134	Navicula flagellijera	1A105	K Y 320290	
Achnanthidium minutissimum	TCC564	KY863481	Navicula gregaria	TA289	KY320297	
$\ A chnanthidium\ pyrenaicum$	TCC832	KY799135	Navicula perminuta	TA413	KY320299	
Achnanthidium reimeri	Arei2	KJ658387	Navicula tripunctata	TCC580	KT072925	
Achnanthidium rivulare	Ariv2	KJ658390	Neidium affine	UTEX FD127	HQ912447	
Achnanthidium straubianum	TCC831	KY799136	Nitzschia aurariae	SZCZCH966	KT943663	
Achnanthidium straubianum	TCC833	KY799137	Nitzschia cf. dubiiformis	SZCZCH970	KT943666	
Aneumastus albanicus	voucher 24426	MH756650	Nitzschia filiformis	UTEX FD267	HQ912453	
Aneumastus stroesei	voucher 23685	MH756663	Nitzschia frustulum	CCMP558	EF423498	
Aneumastus subapiculatus	voucher 24434	MH756665	Nitzschia longissima	KSA2015-9 Nitz.longi-ED	MH064112	
Astartiella sp.	SZCZCH151	KT943613	Parlibellus berkeleyi	SEH015	JX905679	
Auricula complexa	26vi08.1 J.1	KX120569	Parlibellus delognei f. ellipticus	TA387	KY320291	
Auricula mirabilis	4vi08.1cA	KX120567	Parlibellus hamulifer	GU44AK-4	K1577903	
Bacillaria paxillifer	UTEX FD468	HQ912491	Parlibellus hamulifer	SantaRosa cor.green "Trachy-1"	KU179122	
Berkeleva hvalina	ECT3614Bhya	KJ577882	Parlibellus harffianus	SZCZCH75	KT943686	
Berkeleva rutilans	ECT3616	HQ912501	Pauliella taeniata	CCMP1115	KJ658391	
Bolidomonas pacifica	p380	AB430698	Pauliella taeniata	C124	FJ002105	
Bolidomonas pacifica	CCMP1866	HQ912421	Phaeodactylum tricornutum	CCMP2561	HQ912420	
Caloneis cf westii	SZCZCH1002	KT943654	Pinnularia hrehissonii	UTEX FD274	HO912468	
Caloneis lewisii	UTEX ED54	HO912444	Pinnularia termitina	UTEX FD484	HQ912465	
Campuladiscus abraus	mycCA2	KY120583	Placonais alamantis	ED/10	K 1011851	
Campylouiscus ciypeus	3637 C10 100	KX120505	Placonais alginansis	LITEY ED/16	HO012471	
Climaconois viddlogo	5057.C10.100	HO012508	Planothidium caputium	D06 112	VV650812	
Cumaconels riddlede	EC13/24	HQ912508	Planethidium capullum	D00_115	K1050612	
Cocconeis ci. cupulifera	SZCZCH002	K1945080	Planotniaium caputium	BU80-5	K 1 050800	
Cocconeis cl. mascarenica	SZCZCH283	K1943679	subantarcticum	D17_002	KY650822	
Cocconeis cf. sigillata	SZCZCH1200	MT015687	Planothidium cryptolanceolatum	D108_021	KY650818	
Cocconeis pediculus	SpC01	KM084991	Planothidium cryptolanceolatum	D21_002	KY650823	
Cocconeis pediculus	D36_020	KM084977	Planothidium cryptolanceolatum	D26_014	KY650826	

### Table 4 (continued)

Taxon	Strain ID	Assession number ( <i>rbc</i> L)	Taxon	Strain ID	Assession number ( <i>rbc</i> L)
Cocconeis pediculus	Coco1	KM084929	Planothidium frequentissimum	PF1	KJ658392
Cocconeis placentula	UTEX FD23	HQ912456	Planothidium frequentissimum	D06_117b	KY650813
Cocconeis placentula	D36 012	KM084976	Planothidium lanceolatum	PL2	KJ658393
Cocconeis placentula var. euglypta	TCC449	KT072907	Planothidium lanceolatum	PL3	KJ658389
Cocconeis sp.	SZCZP67	KT943614	Planothidium naradoense	D23_024	KY650824
Cocconeis sp.	MPA-2013 ECT3901cocconeid	KC309551	Planothidium sp.	SZCZCH26	KT943678
Cocconeis stauroneiformis	s0230	AB430694	Planothidium suncheonmanense	Ko0408	KY650831
Craspedostauros alvoubii	UTKSA0083	KX981814	Planothidium taeansa	D26 002	KY650825
Craspedostauros amphoroides	CCMP797	KX981815	Prestauroneis integra	AT-177.13	AM710492
Craspedostauros cf. neoconstrictus	CCMP1120	KX981817	Psammodictyon constrictum	GU7X-7 peanut5	KX981830
Craspedostauros paradoxa	GU44BK-1 keeledHcpA25	KX981816	Psammodictyon pustulatum	KSA2015-38 FORAM	MH064134
Craticula accomoda	TCC107	KF959638	Psammothidium abundans	MIC5_40b	KU687468
Craticula ambigua	B060	MH231744	Psammothidium papilio	SHIR_Kbis7	KU687473
Craticula cuspidata	UTEX FD35	HQ912445	Psammothidium subatomoides	B422	KR709269
Craticula cuspidata	Navi4	KM084985	Psammothidium subatomoides	B356	KR709270
Craticula cuspidata	Ind367	MH231745	Psammothidium subatomoides	B352	KR709276
Craticula importuna	AT-70Gel14a	AM710444	Pseudo-nitzschia americana	FBJun06.6	EF423504
Craticula molestiformis	AT-5Nav02	AM710443	Pseudo-nitzschia fraudulenta	BB19	EF423503
Cvlindrotheca closterium	CCMP1855	HQ912509	Rhopalodia contorta	L1299	HQ912392
<i>Cvlindrotheca</i> sp.	UTKSA0079	KX981826	Rhopalodia gibba	CH155	HO912393
<i>Cymatopleura elliptica</i>	L1333	HQ912523	Rossithidium anastasiae	Ros1	KJ658396
Cymatopleura solea	27A	KX120605	Rossithidium petersenii	B354	KR709275
Cymbopleura naviculiformis	22vi092D	KJ011815	Schizostauron davidovichiorum	SZCZP39	KT943604
Cymbopleura sp.	TN-2014 CH254	KJ011817	Schizostauron sp.	SZCZE421	KT943610
Denticula kuetzingii	UTEX FD135	HQ912474	Schizostauron sp.	SZCZE399	KT943609
Dorofeyukea indokotschyi	Ind313	MH231746	Simonsenia aveniformis		KR048205
Dorofeyukea indokotschyi	Ind365	MH231747	Stauroneis acuta	UTEX FD51	HQ912443
Dorofeyukea kotschyi	Ind381	MH231748	Stauroneis cf. anceps	KEL-2015 JAR44_ 3BRUN6	KM999047
Entomoneis pusilla	PMFBIOP1	MF000640	Stauroneis cf. gracilis	KEL-2015 JAR44_ 5ARun6	KM999046
Entomoneis vilicicii	PMFBION4A	MF000637	Stauroneis cf. gracilis	B375	MH231750
Epithemia argus	CH211	HQ912394	Stauroneis gracilis	B540	MH231751
Epithemia turgida	CH154	HQ912396	Stauroneis gracilis	B651	MH231754
Eunotia bilunaris	UTEX FD412	HQ912463	Stauroneis heinii	B644	MH231752
Eunotia glacialis	UTEX FD46	HQ912450	Stauroneis phoenicenteron	AT-182.07	AM710498
Fistulifera pelliculosa	CCMP:543	HQ337547	Stauroneis subgracilis	B376	MH231753
Fistulifera pelliculosa	IK MTA116	JN162792	Stenopterobia curvula	L541	HQ912402
Fistulifera saprophila	TCC535	KF959644	Surirella minuta	UTEX FD320	HQ912522
Fistulifera saprophila	LCR-S-20-3	JQ610165	Surirella splendida	19C	HQ912401
Fistulifera saprophila	TCC508	KC736593	Tryblionella apiculata	UTEX FD465	HQ912464
Fistulifera saprophila	TCC557	KT072923	Tryblionella gaoana	SZCZCH97	KT943683

![](_page_17_Figure_2.jpeg)

Fig. 9 Detail of the position of Cocconeis cf. sigillata MT015687 (SZCZCH1200) in the phylogenetic tree (Fig. 8). See comments on the text

and the major *Cocconeis* clades. Although, the relationship is weakly supported (< 50% ML bootstrap, 0.7763 BI posterior probability).

Finally, *Schizostauron* Grunow, 1867, *Astartiella* Witkowski, Lange-Bertalot & Metzeltin *in* Moser et al. 1998, *Madinithidium* Desrosiers, Witkowski & Riaux-Gobin, *in* Desrosiers et al. 2014, *Karayevia* and *Kolbesia* Round & Bukhtiyarova ex Round, 1998, belong in a monophyletic clade together with Stauroneidaceae as well as *Parlibellus* EJ Cox, 1998 and *Fistulifera* Lange-Bertalot, 1997.

### Discussion

It can be noted that among the morphological characters chosen to delineate groups of taxa in our ultrastructural cladistics analysis (Table 3, Fig. 7), the absence of SV processes (i.e., in *Platessa bahlsii* Potapova, 2012, *P. itoupensis* Tudesque, Le Cohu & CE Wetzel, 2016, *P. kingstonii* Potapova, 2012, *Cocconeis alucitae* Riaux-Gobin & Compère, 2008, *C. neuquina* Frenguelli, 1942, *C. coronatoides* Riaux-Gobin & OE Romero, *in* Riaux-Gobin et al. 2011b) does not tightly group the latter taxa. Thus, it proves

that a more determinant character (than the absence of SV processes), or a combination of characters, better drives the analysis. In the same manner, the terminal raphe endings (simple or showing either more or less large and bent fissures) do not seem determinant to group or individualize taxa.

Following this cladistic analysis, *Cocconeis peltoides* and affiliated taxa (with SV processes), are close to *Psammothidium levanderi* (Hustedt) Bukhtiyarova & Round, 1996 and *P. toroi* S Blanco, Pla-Rabès, CE Wetzel & I Granados, 2017. In contrast, *Cocconeis* taxa without SV processes (such as *C. alucitae*, *C. neuquina* and *C. coronatoides*) are part of another clade, close to several *Platessa* and far from *Achnanthidium*.

Our phylogenetic tree (Fig. 8) suggests that the genus *Cocconeis* is not monophyletic. Surprisingly *C*. cf. *cupulifera* SZCZCH662 (see Riaux-Gobin et al. 2011c, p. 24) has no genetic affiliation to *C*. cf. *sigillata* SZCZCH1200. In

contrast, they both have SV processes and pertain to the same morphological group. Also surprising is the place of C. cf. mascarenica SZCZCH283 (a taxon without SV processes, cf. Riaux-Gobin and Compère 2008) that locates as a sister of C. cf. sigillata (Fig. 8). Several previous investigations (Nakov 2014; Kulikovskiy et al. 2016, 2019; Thomas et al. 2016; Witkowski et al. 2016; Shi et al. 2018) also showed that C. placentula Ehrenberg, 1838 and C. pediculus Ehrenberg, 1838 group together and that C. stauroneiformis Okuno, 1957 pertains to another clade. Finally, and as previously stated by, i.e., Kulikovskiy et al. (2016), the Order Achnanthales arbitrarily groups numerous genera that are not genetically inter-connected, and is thus a polyphyletic group. In more general terms, in support of Thomas et al. (2016), "the monoraphid diatoms are not a natural group." Shape convergence strongly influenced past

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Fig. 10 Cocconeis cf. sigillata MT015687 (SZCZCH1200). a SV in external view, with biseriate striae, *crista marginalis* and SV processes on the mantle; b SV in internal view, with an elliptic SV sternum void of areoalae; c RV in external view; d RV in internal view with two shorter striae between each stria. Striae biseriate near the raphe. Note the Voigt discontinuities (d). Scale bars = 5 µm

![](_page_18_Figure_7.jpeg)

taxonomy, which needs to be revisited in light of genetic approaches. Nevertheless, genetic and morphological approaches (Figs. 7, 10) may complement each other, since the cultivation, followed by sequencing of numerous taxa (i.e., small marine taxa, particularly monoraphid ones) has up to now been impossible (no entries in GenBank), yet essential to clarify their phylogenetic position. Therefore, as a first step, the cladistic analyses based on ultrastructure can help to provide preliminary keys.

**Supplementary Information** The online version contains supplementary material available at https://doi.org/10.1007/s12526-020-01154-9.

Acknowledgments Thanks are due to Chunlian Li for strain isolation (University of Szczecin, Institute of Marine and Environmental Sciences, Szczecin, Poland), Saúl Blanco Lanza (University of León, Department of Biodiversity & Environmental Management), François Féral (University of Perpignan-Via Domitia, Faculty of Law & Economics) and Tamatoa Bambridge (USR3278 CRIOBE EPHE-CNRS-UPVD) for their help with etymology. Peter Esteve and Jeanine Almany (USR3278 CRIOBE EPHE-CNRS-UPVD) are acknowledged, respectively for iconographic support and English revision of the first version of this manuscript, and Yonko Gorand (C2M, University of Perpignan-Via Domitia) for SEM assistance. Yin-chu Wang and Shu-xian Yu are acknowledged for help in sampling campaign performed within Chinese Academy of Sciences President's International Fellowship Initiative (PIFI). We also acknowledge two anonymous reviewers for their helpful comments.

### **Compliance with ethical standards**

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

**Ethical approval** All applicable international, national, and/or institutional guidelines for the care and use of animals were followed by the authors.

**Sampling and field studies** All samplings were done in French Polynesia, under the agreement of CRIOBE (CNRS–USR 3278, France). Diatoms are not protected organisms.

**Data availability** The datasets generated during and/or analyzed during the current study, particularly all SEM images, are available from the corresponding author on reasonable request. No citations of other datasets.

Author contribution CR-G, AW, and EG conceived and designed research, and analyzed data. CR-G conducted the sampling and ultrastructural description of the new taxon. EG and AW conducted the genetic analysis of a close taxon. PS-A conducted the cladistic analysis on ultrastructural characters. LE managed bibliographic research. GD-K conducted the technical assistance. CR-G wrote the manuscript with help of EG and all co-authors. All authors read and approved the manuscript.

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