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OPEN Green armoured tardigrades (Echiniscidae: Viridiscus), including a new species from the Southern Nearctic, exemplify problems with tardigrade variability research

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Ranges of tardigrade intraspecific and interspecific variability are not precisely defined, both in terms of morphology and genetics, rendering descriptions of new taxa a cumbersome task. This contribution enhances the morphological and molecular dataset available for the heterotardigrade genus Viridiscus by supplying new information on Southern Nearctic populations of V. perviridis, V. viridianus, and a new species from Tennessee. We demonstrate that, putting aside already well-documented cases of significant variability in chaetotaxy, the dorsal plate sculpturing and other useful diagnostic characters, such as morphology of clavae and pedal platelets, may also be more phenotypically plastic characters at the species level than previously assumed. As a result of our integrative analyses, V. viridianus is redescribed, V. celatus sp. nov. described, and V. clavispinosus designated as nomen inquirendum, and its junior synonymy with regard to V. viridianus suggested. Morphs of three Viridiscus species (V. perviridis, V. viridianus, and V. viridissimus) are depicted, and the implications for general echiniscid taxonomy are drawn. We emphasise that taxonomic conclusions reached solely through morphological or molecular analyses lead to a distorted view on tardigrade α -diversity.

Tardigrades, also known as water bears or moss piglets, represent an invertebrate phylum closely related to onychophorans and arthropods within Panarthropoda¹, which consists of *ca*. 1500 described species to date². These are grouped into two classes: Heterotardigrada and Eutardigrada. Heterotardigrades are characterised by the presence of cirrus A^{3-5} , whereas eutardigrades are generally distinguishable by their simplified, vermiform external morphology⁶. Tardigrades inhabit limno-terrestrial and aquatic, both freshwater and marine, habitats, the former typically in association with bryophytes, algae, lichens, or leaf litter^{5,7,8}. Although the number of documented tardigrade species has increased significantly in the past three decades, most regions globally remain sparsely sampled for these diminutive animals^{9,10}. The distributions of a few terrestrial tardigrade species are well documented for some regions of North America, e.g. see11, but the general paucity of North American tardigrade specialists has left much of the continent unexplored in this regard (most recently reviewed in^{12,13}).

The Echiniscidae, the most speciose family within the class Heterotardigrada¹⁴, contain the morphologically odd genus Viridiscus¹⁵⁻¹⁷. The former Echiniscus viridis group was established by Ramazzotti¹⁸, who considered E. viridis Murray, 1910¹⁹, E. perviridis Ramazzotti, 1959¹⁸, E. viridissimus Péterfi, 1956²⁰, and E. rufoviridis du Bois-Reymond Marcus, 1944²¹ to be closely related based on green cuticular pigments in plates forming the armour, alongside orange pigments typically present in the body cavity of echiniscids²². Two further species were

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added to this group many decades later: *E. viridianus* Pilato et al., 2007²³ and *E. clavispinosus* Fontoura et al., 2011²⁴. *Viridiscus* was erected based on morphological and molecular data to accommodate the distinctiveness of this group: dark body pigments, well-developed, sabre-like claws, a lack of trunk appendages except cirrus *A*, and typical dorsal plate sculpture composed of an intracuticular sponge layer and small, flat, densely arranged epicuticular granules¹⁵. Recently, new evidence was presented, pointing out possibly large intraspecific variability within *Viridiscus*, as two morphs were distinguished within *V. viridissimus*^{16,17}. Moreover, *V. rufoviridis* was transferred to *Barbaria*²⁵, rectifying the earlier assignment of this species within the *viridis* group^{18,23,26}. Consequently, *Viridiscus* currently comprises five species: *V. clavispinosus*, *V. perviridis*, *V. viridianus*, *V. viridis*, and *V. viridissimus*.

Here, we describe a new species of *Viridiscus* found in previously described *Viridiscus* assemblages in Tennessee^{16,27} and identified as *V*. aff. *viridianus* in¹⁷. We also redescribe *V. viridianus* based on multiple populations collected in Alabama and Florida. Newly obtained data confirm and strengthen previous findings that *Viridiscus* species are variable in the dorsal plate sculpturing¹⁷ and morphology of the primary clavae. We update the genus phylogeny, and question the validity of *V. clavispinosus*, which we hypothesise to be a synonym of *V. viridianus*. We stress that there is no convincing evidence for the presence of *V. viridis* outside of the Hawaiian Archipelago²⁶, and the Nearctic records²⁸ of this species are unreliable. The case of *Viridiscus* variability yet again illustrates that forming taxonomic conclusions based solely on morphology is unadvised and should be renounced¹⁷, especially since DNA sequence-based tools are readily available to incorporate into the systematics of these organisms.

Results

General remarks: morphology and phylogeny

There are two general morphotypes distinguishable within *Viridiscus* with respect to the dorsal plate sculpturing: (I) pore-dominated, in which epicuticular granules are restricted to anterior portions of paired segmental plates (a usual morph of *V. viridissimus*, see Fig. 1A; however, we found several specimens with developmental aberrations embracing fusion of different elements of dorsal armour, see Fig. 1B); and (II) granule-dominated, in which pores are absent (the remaining four species, represented herein by *V. viridianus*: Fig. 1C). This division is also clearly visible in SEM: pores of *V. viridissimus* (Fig. 2A,B) are approaching the size of epicuticular granules of *V. perviridis* (Fig. 2C,D). However, the SEM analysis revealed also another element of sculpturing in the second *Viridiscus* morphotype: micropores, $\ll 1$ µm in diameter and distributed irregularly between the granules (Fig. 2C,D).

Viridiscus viridissimus exhibits considerable intra-specific variation in the morphology of dorsal armour¹⁷. In the typical morph (Fig. 3A), reliably recorded from the Holarctic and Oriental regions¹⁷, epicuticular granules are limited to the anterior portion of the second median and paired segmental plates, and to the third median plate. In contrast, the much less common morph of V. viridissimus (initially described as V. miraviridis Nelson et al., 2020¹⁶), so far identified only among moss samples from Tennessee, exhibits epicuticular granules in all plate portions (Fig. 3B). Having the possibility to analyse abundant populations of V. perviridis and V. viridianus, we discovered atypical morphs in these species, too. In V. perviridis, the usually well-discernible epicuticular granules (Fig. 3C) can be poorly developed (and therefore blurred with endocuticle in PCM), especially in the scapular and caudal (terminal) plates (Fig. 3D). In V. viridianus, the reduction of epicuticular granules (Fig. 3E) can be even more pronounced to the extent that granules are absent, and only the intracuticular sponge layer is identifiable in PCM (Fig. 3F). These atypical morphs were not associated with a particular life stage or sex but appeared in large monospecific populations of the analysed Viridiscus spp. No atypical morphs were observed in the case of V. celatus sp. nov. (Fig. 3G,H), but the available sample size was significantly smaller in the case of the new species compared to the other analysed Viridiscus spp. As already stated in the diagnosis of the genus¹⁵, all known larvae of Viridiscus spp., irrespectively of the adult morphotype (type I or II), possess large pores beside granules (Fig. 4). We confirmed the presence of such pores in larvae of V. perviridis, V. viridianus, and V. viridissimus, and they have been also detected in V. clavispinosus²⁴. Larvae of V. viridis s.s. (inhabiting the Hawaiian Archipelago) have never been found^{19,26}.

The phylogeny fully conformed with the morphological analyses, indicating the presence of four species (Fig. 5), relationships among which were as follows: (((*V. perviridis* (*V. celatus* **sp. nov.** (*V. viridianus* + *V. viridissimus*))). *Viridiscus celatus* **sp. nov.** was classified in¹⁷ as *V.* aff. *viridianus*. Importantly, species delimitations based on COI did not distinguish between *V. perviridis* and *V. celatus* **sp. nov.**, which are clearly separated based on both ITS markers (Fig. 5) and morphology, pinpointing the questionable utility of the COI marker in tardigrade species delimitation. Given the abundant material and the proximity of the type locality in Auburn (Alabama) to the localities sampled in this study, we redescribe *V. viridianus* to provide a detailed insight into its intraspecific variability.

Integrative redescription of Viridiscus viridianus (Pilato et al., 2007)

(Tables 1, 2, 3, 4, Figs. 6, 7, 8, 9, 10, 11, raw morphometry in Supplementary Material 1).

New material examined. Populations from Alabama and Florida, 346 specimens in total were processed for PCM, SEM and DNA analyses (Table 5); additional few hundred specimens were frozen for future analyses. Type locality. North America, USA, Alabama, Auburn.

Additional localities. North America, USA, New Mexico²³; the Azores, Ribeira Fria, Lages do Pico^{23,29,30}; North America, USA, New Jersey²⁸; Central America, the Lesser Antilles, Antigua²⁸. Given the reported variability in the pattern of the dorsal armour, these additional localities should be verified.

Etymology. From Latin *viridianus* = greenish. An adjective in nominative singular.



Figure 1. Two main morphotypes present in *Viridiscus* (PCM): (**A**) pores present, epicuticular granules typically reduced: the usual morph of *V. viridissimus* (Tennessee, dorsal view); (**B**) an aberrant specimen of *V. viridissimus* (Tennessee, dorsolateral view); (**B**) an aberrant specimen of *V. viridissimus* (Tennessee, dorsolateral view); (**B**) an aberrant specimen of *V. viridissimus* (Tennessee, dorsolateral view); (**B**) an aberrant specimen of *V. viridissimus* (Tennessee, dorsolateral view); (**B**) an aberrant specimen of *V. viridissimus* (Tennessee, dorsolateral view); (**B**) an aberrant specimen of *V. viridissimus* (Tennessee, dorsolateral view); (**B**) an aberrant specimen of *V. viridissimus* (Tennessee, dorsolateral view); (**C**) epicuticular granules dominant, pores present only in larval stage: *V. viridianus* (Alabama, dorsolateral view). Scale bars = 50 µm.



Figure 2. Two main morphotypes of dorsal plate sculpturing present in *Viridiscus* (SEM): pores dominant, only *V. viridissimus* (the population from Vietnam): (**A**) a fragment of the scapular plate; (**B**) close up of the posterior portion of the second paired segmental plate; epicuticular granules dominant, micropores visible only in SEM, all remaining *Viridiscus* species (the population of *V. perviridis* from Vietnam shown): (**C**) a fragment of the scapular plate; (**D**) close up of the central portion of the scapular plate. Scale bars in µm.

Animals. Females (i.e., from the third instar onwards; measurements and statistics in Table 1). Body cavity with yellowish pigments (typical for most echiniscids), whereas dorsal and pedal cuticular plates light to dark green (Figs. 1C, 3E,F). Red eves and yellow pigments present in live specimens, but dissolve after mounting in Hoyer's medium, thus only green pigmentation persists. Body bulky (Fig. 6), with a poorly delimited cephalic region (Fig. 7). The cervical (neck) plate is well-developed, but sculptureless (Fig. 7). Weakly developed lateralmost, rectangular portions of the scapular plate with a weak sculpturing (Fig. 1C). Dorsal plate sculpturing ordinarily comprising polygonal epicuticular granules with scarce micropores, barely identifiable, even with SEM (Fig. 8A). Lateral and ventral endocuticle with intracuticular pillars, visible in PCM as minute dark dots, but identifiable in SEM only when the thin epicuticle is ruptured (Fig. 8B-D). Pillars larger and more sclerotised in proximal and central limb portions, forming longitudinal, narrow pulvini, and pedal platelets, respectively (Fig. 9). Some specimens exhibit a differently formed central pedal portion, more convex than the remainder of each platelet (Figs. 9, 11E). Areas of more sclerotised pillars always form a pair of merged subcephalic plates (Fig. 10F). Cirrus A short (<20% of body length) and thin. A remarkable diversity of primary clava shapes: some specimens within a population have both clavae pointed and clearly conoid (Figs. 7A, 10A-C), some have both clavae dactyloid, i.e., elongated, but without a pointed tip (Fig. 10D), and in some individuals both clavae are tubby, i.e., of a typical, Echiniscus-type shape (Fig. 10E). Importantly, numerous specimens showed a mixture of these shapes, that is the clava of one specimen differed in morphology from the other one on the same specimen. Claws massive and isonych (Figs. 10G-I, 11A-D).

Males (i.e., most probably from the third instar onwards; measurements and statistics in Table 2). No detectable sexual dimorphism besides the circular gonopore.

Juveniles (i.e., the second instar; measurements and statistics in Table 3). Smaller than adults, but qualitatively like them. Gonopore absent.

Larvae (i.e., the first instar; measurements and statistics in Table 4). Body size overlaps with juveniles. Anterior portions of paired segmental plates, and median plate 2 sculptureless. Large cuticular pores in the dorsal armour. No gonopore or anus.

Eggs. Up to five orange eggs per shed exuvia, but typically fewer (see³¹).

Remarks. Males were present in all examined populations of the species.



Figure 3. Intrageneric and intraspecific variability in *Viridiscus* (PCM): (**A**) *V. viridissimus* (Tennessee), the typical morph; (**B**) *V. viridissimus* (Tennessee), the atypical morph with well-developed epicuticular granules¹⁶; (**C**) *V. perviridis* (Madeira), the typical morph with well-developed epicuticular granules; (**D**) *V. perviridis* (Alabama), the atypical morph with poorly delineated epicuticular granules; (**E**) *V. viridianus* (Alabama), the typical morph with well-developed epicuticular granules; (**F**) *V. viridianus* (Alabama), the typical morph with epicuticular granules absent (arrowhead points out conoid primary clava); (**G**,**H**) *V. celatus* sp. nov. (Tennessee), the only morph observed. Scale bars = 20 µm.

Description of Viridiscus celatus sp. nov. Momeni, Gąsiorek, Nelson & Michalczyk

(Tables 6, 7, Figs. 12, 13, 14, raw morphometry in Supplementary Material 2). ZooBank registration number: urn:lsid:zoobank.org:act:7E5416A6-E49F-46C2-8BA6-3D734B2A10A4.

Material examined. Populations from Tennessee, 19 specimens in total processed for PCM and DNA analyses (Table 5).

Type locality. 36°18′N, 82°22′W, *ca.* 520 m asl: USA, Tennessee, Washington County, Johnson City. *Grimmia* sp. mosses from a concrete cap on a brick fence post.

Etymology. From Latin *celatus* = hidden, concealed. The name refers to the fact that the new species was not identified as new taxon for a long time, although the locality has been extensively sampled for tardigrades (16,27 referred to the species as *V. perviridis* based on the identification by Maucci³², and morphological characters, including cirrus *A* length). An adjective in nominative singular.



Figure 4. Larva of V. viridianus (PCM, dorsolateral view). Scale bar = 20 µm.



Figure 5. Phylogenetic relationships of the genus *Viridiscus*: Bayesian tree based on the concatenated ITS-1 + ITS-2 dataset (1058 bp); vertical bars denote different delineation methods used in the formulation of the primary molecular species hypotheses: ASAP, ABGD, and bPTP; (COI) refers to COI delimitation in all three methods). Asterisks indicate the maximal (1.00/100) posterior probability/bootstrap value. *Echiniscus succineus* was used as an outgroup. Scale bar represents substitutions per position.

Type depositories: Type series: holotypic female (slide US.081.03), allotypic male (slide US.081.02), and nine paratypes (slides US.078.03 and US.081.03-4), are deposited at the Faculty of Biology, Jagiellonian University (Kraków, Poland).

Animals. Females (i.e., from the third instar onwards; measurements and statistics in Table 6). Body mediumsized and bulky. Body cavity with yellowish pigments (typical for most echiniscids), whereas dorsal and pedal cuticular plates olive green (Figs. 1C, 3E,F). Red eyes and yellow pigments present in live specimens, but dissolve after mounting in Hoyer's medium, thus only green pigmentation persists (Figs. 12, 13B). Except for cirrus *A*, with a tubby clava near the cirrophore (Fig. 12), other dorsal and lateral trunk appendages are absent. Cephalic

| | | Range | | Mean | | SD | |
|-------------------------------|----|-----------|-----------|------|------|-----|-----|
| Character | N | μm | sp | μm | sp | μm | sp |
| Body length | 15 | 220-326 | 457-573 | 280 | 527 | 29 | 29 |
| Scapular plate length | 15 | 44.3-57.9 | - | 53.0 | - | 3.6 | - |
| Head appendage lengths | | | | | | | |
| Cirrus internus | 14 | 9.0-16.7 | 16.8-28.8 | 12.5 | 23.5 | 2.0 | 2.9 |
| Cephalic papilla | 15 | 6.4-8.9 | 12.4–16.3 | 7.9 | 14.9 | 0.7 | 1.0 |
| Cirrus externus | 15 | 14.0-24.6 | 28.2-42.5 | 18.7 | 35.2 | 2.9 | 3.9 |
| Clava | 15 | 5.4-7.6 | 10.3-13.3 | 6.4 | 12.1 | 0.7 | 0.9 |
| Cirrus A | 15 | 24.5-46.1 | 53.0-79.9 | 37.7 | 70.9 | 5.9 | 8.0 |
| Cirrus A/body length ratio | 15 | 9-15% | - | 13% | - | 2% | - |
| Body appendage lengths | | | | | | | |
| Spine on leg I length | 14 | 2.2-3.6 | 4.3-6.9 | 2.7 | 5.1 | 0.4 | 0.7 |
| Papilla on leg IV length | 15 | 3.0-4.6 | 6.7-8.6 | 3.9 | 7.3 | 0.4 | 0.5 |
| Number of teeth on the collar | 15 | 9–12 | - | 10.9 | - | 1.1 | - |
| Claw I heights | | | | | | | |
| Branch | 15 | 15.8-19.7 | 28.5-36.6 | 17.4 | 32.9 | 1.2 | 2.1 |
| Spur | 13 | 1.9-2.5 | 3.3-5.2 | 2.2 | 4.1 | 0.2 | 0.5 |
| Spur/branch height ratio | 13 | 10-14% | - | 13% | - | 1% | - |
| Claw II heights | • | | | | | | |
| Branch | 15 | 14.9-19.0 | 30.9-33.6 | 17.0 | 32.0 | 1.0 | 0.8 |
| Spur | 15 | 1.9-2.5 | 3.5-5.4 | 2.2 | 4.1 | 0.2 | 0.4 |
| Spur/branch height ratio | 15 | 11-16% | - | 13% | - | 1% | - |
| Claw III heights | | | | | | | |
| Branch | 15 | 14.7-19.0 | 29.8-33.2 | 16.8 | 31.8 | 1.2 | 1.1 |
| Spur | 14 | 1.7-2.4 | 3.2-4.7 | 2.1 | 3.9 | 0.2 | 0.4 |
| Spur/branch height ratio | 14 | 10-15% | - | 12% | - | 1% | - |
| Claw IV heights | | | - | | - | | |
| Branch | 15 | 17.3-20.9 | 33.7-39.1 | 19.3 | 36.4 | 1.1 | 1.3 |
| Spur | 11 | 2.4-3.5 | 4.4-6.3 | 2.9 | 5.3 | 0.4 | 0.5 |
| Spur/branch height ratio | 11 | 12-17% | - | 15% | - | 2% | - |

Table 1. Measurements (in μ m) of selected morphological structures of adult females of *Viridiscus viridianus* mounted in Hoyer's medium. *sp* the proportion between the length of a given structure and the length of the scapular plate, ? unknown.

appendages include internal and external peribuccal cirri with tubby cephalic papillae between them (Fig. 12). Dorsal plate sculpturing comprises large epicuticular granules (Fig. 12), which may be poorly developed in central plate portions (Fig. 13B). Sponge layer identifiable beneath granules. Granules appear more convex in anterior portions of paired segmental plates than in the remainder of the armour in PCM. Micropores not visible in PCM and their presence or absence remains to be confirmed in SEM.

All plates strongly sclerotised and with clear edges. The cephalic plate with a well-marked anterior chaliceshaped incision, the cervical plate and lateral sections of the body lack dense granulation and are covered with fine regular punctuation. The scapular plate contains three portions. Only the central part is visible in the dorsal view, and two small, weakly delineated, trapezoidal sections are present on the lateral portions of the body, with intracuticular pillars visible (Fig. 13B). The first median plate is triangular and unipartite, the second median plate is subdivided into two portions, and the anterior portion lacks the sponge layer. The third median plate is absent, but the area between the paired segmental plate II and the caudal plate is covered with large granules. Paired segmental plates I and II have two clearly delineated parts. Intersegmental plate is inserted between the posterolateral edge of the paired segmental plate I and anterior margin of paired segmental plate II. The caudal incisions are unsclerotised and weakly marked (Figs. 12, 13B).

Venter densely granulated in PCM (endocuticular pillars); a pair of subcephalic plates present (Fig. 14A). Gonopore hexapartite. Pulvini (= narrow proximal bands of intracuticular pillars) and pedal platelets (= broad central bands of pillars) are visible on all legs. Dentate collar with numerous irregular teeth (Fig. 14C). Sensory organs present on all legs: a tiny spine on leg I embedded at the edge of pedal platelet; hemispherical rudimentary papillae on legs II–III, embedded in the centre of pedal platelets (identifiable only when specimens are dorso-laterally oriented); and papilla IV on hind legs (Fig. 13B). Claws anisonych; primary spurs I–III tiny and thin, positioned slightly lower on branches compared to more massive spurs IV (Fig. 14B–C).

Males (i.e., most probably from the third instar onwards; measurements and statistics in Table 7). No sexual dimorphism observable in body size or qualitative traits (Fig. 13A). Gonopore circular.

| | | Range | | Mean | | SD | |
|-------------------------------|---|-----------|-----------|------|------|-----|------|
| Character | N | μm | sp | μm | sp | μm | sp |
| Body length | 4 | 248-257 | 521-579 | 254 | 546 | 4 | 24 |
| Scapular plate length | 4 | 42.8-48.9 | - | 46.6 | - | 2.6 | - |
| Head appendage lengths | | | | | | | |
| Cirrus internus | 4 | 11.9-13.4 | 26.2-28.3 | 12.7 | 27.2 | 0.7 | 1.0 |
| Cephalic papilla | 4 | 6.7-8.1 | 15.7–17.1 | 7.7 | 16.4 | 0.6 | 0.6 |
| Cirrus externus | 4 | 14.1-19.3 | 32.9-40.7 | 17.1 | 36.5 | 2.3 | 3.4 |
| Clava | 4 | 5.4-7.1 | 12.6-15.0 | 6.5 | 13.9 | 0.8 | 1.1 |
| Cirrus A | 3 | 25.2-37.1 | 58.9-78.3 | 33.1 | 71.7 | 6.8 | 11.1 |
| Cirrus A/body length ratio | 3 | 10-14% | - | 13% | - | 2% | - |
| Body appendage lengths | | | | | | | |
| Spine on leg I length | 4 | 1.5-3.0 | 3.5-6.3 | 2.5 | 5.4 | 0.7 | 1.3 |
| Papilla on leg IV length | 4 | 3.0-4.0 | 7.0-8.4 | 3.7 | 7.9 | 0.5 | 0.6 |
| Number of teeth on the collar | 4 | 9-14 | - | 10.5 | - | 2.4 | - |
| Claw I heights | | | | | | | |
| Branch | 4 | 15.2-17.1 | 33.8-35.5 | 16.1 | 34.6 | 0.8 | 0.8 |
| Spur | 3 | 1.7-2.2 | 3.5-5.1 | 2.0 | 4.4 | 0.3 | 0.9 |
| Spur/branch height ratio | 3 | 10-14% | - | 13% | - | 2% | - |
| Claw II heights | | | | | | | |
| Branch | 4 | 14.5-16.2 | 32.1-33.9 | 15.3 | 32.9 | 0.7 | 0.8 |
| Spur | 4 | 1.8-2.3 | 4.2-4.9 | 2.1 | 4.4 | 0.2 | 0.3 |
| Spur/branch height ratio | 4 | 12-15% | - | 13% | - | 1% | - |
| Claw III heights | | | | | | | |
| Branch | 4 | 13.5-16.2 | 31.5-33.1 | 15.1 | 32.4 | 1.1 | 0.7 |
| Spur | 4 | 1.8-2.2 | 3.7-4.7 | 2.0 | 4.3 | 0.2 | 0.5 |
| Spur/branch height ratio | 4 | 11-15% | - | 13% | - | 2% | - |
| Claw IV heights | - | | | - | | | |
| Branch | 4 | 17.3-19.4 | 36.6-40.4 | 18.0 | 38.7 | 1.0 | 1.7 |
| Spur | 3 | 2.7-3.1 | 5.9-6.5 | 2.9 | 6.3 | 0.2 | 0.3 |
| Spur/branch height ratio | 3 | 16-17% | - | 16% | - | 1% | - |

Table 2. Measurements (in μ m) of selected morphological structures of adult males of *Viridiscus viridianus* mounted in Hoyer's medium. *sp* the proportion between the length of a given structure and the length of the scapular plate, ? unknown.

Juveniles (i.e., the second instar; measurements and statistics in Table 7). Smaller than both females and males. Qualitatively like adults; gonopore absent.

- Larvae. Not found.
- Eggs. Not found.

Remarks. Found only in association with large populations of V. viridissimus¹⁷.

Differential diagnosis. The new species from Tennessee is differentiated from all *Viridiscus* spp. based on the presence of plesiomorphic papillae on legs II–III. These structures are, however, barely identifiable in specimens oriented dorsoventrally, hence we enumerate other criteria making *V. celatus* **sp. nov.** distinct from:

- *Viridiscus clavispinosus*, by the relative length of cirrus *A* (34–49% vs <15% of the body length), and primary spurs IV less divergent from claw branches;
- Viridiscus perviridis, veritably reported from the Holarctic and Oriental regions¹⁷, by the length of cirrus A (34–49% vs typically \gg 50% of the body length, see the subsection below that addresses this character), the weakly developed caudal incisions (strongly sclerotised and well-marked in all syntypes of *V. perviridis*), and the body colour (light to olive green vs usually dark green to almost black in *V. perviridis*, also in mounted specimens);
- *Viridiscus viridianus*, reliably reported only from the USA, by the relative length of cirrus *A* (34–49% vs <20% of the body length), the lack of pedal platelets with a distinctly formed central portion, and the more pronounced sculpturing of the anterior portions of paired segmental plates;
- *Viridiscus viridis*, reliably reported only from the Hawaiian Archipelago^{19,26}, by the relative length of cirrus A (34–49% vs < 10% of the body length), and a different pattern of dorsal sculpturing (in general *V. viridis* has noticeably fewer epicuticular granules on all plates, see fig. 1 in²⁶);
- *Viridiscus viridissimus*, with a likely wide distribution in the Holarctic, Oriental, and Neotropical regions^{17,33}, by the absence of pores in dorsal armour, and a better developed sponge layer of cuticle.

| | | Range | | Mean | | SD | |
|-------------------------------|---|-----------|-----------|------|------|-----|-----|
| Character | N | μm | sp | μm | sp | μm | sp |
| Body length | 7 | 124-202 | 419–582 | 171 | 503 | 26 | 49 |
| Scapular plate length | 7 | 29.6-39.2 | - | 33.9 | - | 3.2 | - |
| Head appendage lengths | | | | | | | |
| Cirrus internus | 6 | 6.9-9.2 | 20.2-25.6 | 7.9 | 23.2 | 1.0 | 2.0 |
| Cephalic papilla | 7 | 5.0-6.2 | 14.3–18.1 | 5.5 | 16.2 | 0.4 | 1.4 |
| Cirrus externus | 7 | 7.8-14.4 | 25.1-36.7 | 10.1 | 29.5 | 2.2 | 3.6 |
| Clava | 6 | 4.1-5.3 | 11.5–15.5 | 4.5 | 13.1 | 0.4 | 1.5 |
| Cirrus A | 5 | 20.2-26.9 | 58.9-72.6 | 22.5 | 65.0 | 2.7 | 5.7 |
| Cirrus A/body length ratio | 5 | 12-17% | - | 13% | - | 2% | - |
| Body appendage lengths | | | | | | | |
| Spine on leg I length | 5 | 1.4-1.7 | 4.0-4.6 | 1.5 | 4.4 | 0.1 | 0.2 |
| Papilla on leg IV length | 7 | 2.0-2.8 | 6.4-8.4 | 2.5 | 7.3 | 0.3 | 0.6 |
| Number of teeth on the collar | 6 | 6-11 | - | 8.3 | - | 1.8 | - |
| Claw I heights | | | | | | | |
| Branch | 7 | 9.4-13.2 | 30.9-33.7 | 10.9 | 32.1 | 1.3 | 1.0 |
| Spur | 7 | 1.4-2.0 | 4.0-5.8 | 1.7 | 5.1 | 0.2 | 0.6 |
| Spur/branch height ratio | 7 | 12-19% | - | 16% | - | 2% | - |
| Claw II heights | | | | | | | |
| Branch | 7 | 9.0-13.0 | 30.0-33.2 | 10.5 | 31.1 | 1.3 | 1.2 |
| Spur | 7 | 1.3-1.9 | 4.0-5.9 | 1.7 | 4.9 | 0.2 | 0.6 |
| Spur/branch height ratio | 7 | 13-18% | - | 16% | - | 2% | - |
| Claw III heights | | | | | | | |
| Branch | 7 | 8.4-12.2 | 28.3-31.1 | 10.1 | 29.7 | 1.3 | 1.1 |
| Spur | 6 | 1.4-1.8 | 4.0-4.7 | 1.6 | 4.5 | 0.2 | 0.2 |
| Spur/branch height ratio | 6 | 13-16% | - | 15% | - | 1% | - |
| Claw IV heights | | | | | | | |
| Branch | 7 | 10.1-14.3 | 32.4-36.8 | 11.9 | 34.9 | 1.5 | 1.7 |
| Spur | 7 | 1.7-2.3 | 5.2-7.1 | 2.0 | 6.0 | 0.2 | 0.7 |
| Spur/branch height ratio | 7 | 14-20% | - | 17% | - | 2% | - |

Table 3. Measurements (in μ m) of selected morphological structures of juveniles of *Viridiscus viridianus* mounted in Hoyer's medium. *sp* the proportion between the length of a given structure and the length of the scapular plate, ? unknown.

Discussion

Viridiscus viridianus vs Viridiscus clavispinosus

Viridiscus viridianus was described by Pilato et al.²³ based on populations from Alabama (the origin of the holotype, thus locus typicus), New Mexico (the Nearctic region), and the Azores (Macaronesia, the westernmost part of the Palaearctic region). A few years later, Fontoura et al.²⁴ described V. clavispinosus (from the Archipelago of Cape Verde, which is also a part of Macaronesia, but located southwards of the Azores), using a seemingly sound autapomorphy, namely conoid primary clavae (with pointed apices). However, the data gathered in the present study undermine the validity of V. clavispinosus, as the shape of primary clava is evidently a variable character in some species of *Viridiscus* (Fig. 10A–E), rendering it unsuitable for species delineation. The surface between the paired segmental plate II and the caudal plate can be weakly sculptured (Figs. 1C, 3E, 6A,C) or unsculptured (Fig. 3F) in a population of one species. Contrary to what was stated in²⁴, neither V. viridianus nor V. clavispinosus exhibit well-marked epicuticular granules in the cephalic and cervical plates (compare figs. 1b, e in²³, fig. 1a in²⁴, and Fig. 7 herein). In echiniscid species, the dorsal plate sculpturing is less pronounced in these two anteriormost elements of armour. The sculpturing of the less sclerotised anterior portion of median plate 2 can also be more or less developed, and sometimes absent within a single population. The remaining characters referring to the dorsal sculpturing mentioned in²⁴ also are invalid in light of various atypical morphs presented for *Viridiscus* species in this paper, including V. viridianus. We refrained however, from synonymising V. clavispinosus with V. viridianus for two reasons: (1) as rightly stated by Fontoura et al.²⁴, primary spurs IV of V. clavispinosus are more developed than those of V. viridianus (compare fig. 2c in²⁴ and Fig. 11D herein); (2) although V. viridianus potentially has a wide circum-Atlantic distribution (whilst molecular evidence is lacking), V. clavispinosus is known from a different part of Macaronesia. Therefore, V. clavispinosus is designated as nomen inquirendum, given the substantial doubts regarding its separateness from V. viridianus described above.

| | | Range | | Mean | | SD | |
|-------------------------------|---|-----------|-----------|------|------|-----|-----|
| Character | N | μm | sp | μm | sp | μm | sp |
| Body length | 7 | 116-147 | 431-535 | 134 | 504 | 13 | 44 |
| Scapular plate length | 7 | 25.7-27.5 | - | 26.6 | - | 0.8 | - |
| Head appendage lengths | | | | | | | |
| Cirrus internus | 6 | 5.2-6.8 | 19.0-25.1 | 5.8 | 21.5 | 0.6 | 2.1 |
| Cephalic papilla | 6 | 4.4-4.9 | 16.2–18.2 | 4.7 | 17.5 | 0.2 | 0.7 |
| Cirrus externus | 7 | 6.1-8.9 | 23.4-33.1 | 7.4 | 27.7 | 0.9 | 3.1 |
| Clava | 6 | 3.5-4.7 | 13.4–17.3 | 4.2 | 15.7 | 0.4 | 1.5 |
| Cirrus A | 6 | 13.8-17.5 | 53.5-64.6 | 15.7 | 58.5 | 1.6 | 5.0 |
| Cirrus A/body length ratio | 6 | 10-14% | - | 11% | - | 2% | - |
| Body appendage lengths | | | | | | | |
| Spine on leg I length | 7 | 1.3-1.5 | 4.7-5.8 | 1.4 | 5.3 | 0.1 | 0.4 |
| Papilla on leg IV length | 6 | 2.0-2.3 | 7.3-8.6 | 2.2 | 8.1 | 0.1 | 0.5 |
| Number of teeth on the collar | 7 | 6–9 | - | 8.0 | - | 1.2 | - |
| Claw I heights | | | | | | | |
| Branch | 7 | 9.1-10.1 | 35.4-37.6 | 9.7 | 36.5 | 0.3 | 0.9 |
| Spur | 7 | 1.7-2.4 | 6.6-8.8 | 2.0 | 7.5 | 0.3 | 0.9 |
| Spur/branch height ratio | 7 | 19-25% | - | 21% | - | 2% | - |
| Claw II heights | | | | | | | |
| Branch | 7 | 8.7-10.0 | 33.3-36.4 | 9.4 | 35.1 | 0.5 | 1.2 |
| Spur | 7 | 1.7-2.2 | 6.6-8.1 | 1.9 | 7.3 | 0.2 | 0.6 |
| Spur/branch height ratio | 7 | 19-24% | - | 21% | - | 2% | - |
| Claw III heights | | | | | | | |
| Branch | 7 | 8.4-9.7 | 32.7-35.8 | 9.1 | 34.0 | 0.5 | 1.2 |
| Spur | 5 | 1.7-2.2 | 6.3-8.2 | 1.9 | 7.1 | 0.2 | 0.7 |
| Spur/branch height ratio | 5 | 18-24% | - | 21% | - | 3% | - |
| Claw IV heights | | | | | | | |
| Branch | 7 | 9.8-10.8 | 37.8-40.1 | 10.3 | 38.7 | 0.3 | 0.9 |
| Spur | 4 | 2.1-2.4 | 7.7-8.9 | 2.3 | 8.5 | 0.1 | 0.6 |
| Spur/branch height ratio | 4 | 19-23% | - | 22% | - | 2% | - |

Table 4. Measurements (in μ m) of selected morphological structures of larvae of *Viridiscus viridianus* mounted in Hoyer's medium. *sp* the proportion between the length of a given structure and the length of the scapular plate, ? unknown.

Inter- vs intraspecific variability in Viridiscus and its consequences for echiniscid taxonomy

One of the prime tasks for taxonomists is recognising the borders of inter- and intraspecific variability in a given group of organisms (for microscopic animals, see e.g.³⁴⁻³⁷). In the preliminary study that integratively addressed variability of Viridiscus¹⁷, we accentuated the role of combining molecular and morphological approaches in tardigrade taxonomy, as utilising only one line of evidence either leads to taxonomic inflation (e.g., the synonymy of V. miraviridis^{16,17}) or deflation (e.g., the Milnesium case³⁸). We have already underscored high variability of characters previously deemed as universally stable at the species level in the Echiniscidae (cirrus A length, details of cuticular ornamentation) in Echiniscus and Nebularmis spp.^{39,40}. This study adds further observations on atypical morphs within Viridiscus, which show disparities with typical morphs of a given species that are larger than interspecific variability in multiple new characters, such as the general phenotype of the dorsal sculpturing, the shape of primary clava, and morphology of pedal platelets. It is gradually becoming obvious that even within such a morphologically coherent family as Echiniscidae⁴¹, phenotypic plasticity of diagnostic characters varies between lineages at various taxonomic levels (genus, species, etc.), as it happens with the shape of primary clavae in Viridiscus, which is labile in contrast to other known echiniscid genera. Consequently, conservative characters in one genus should not simply be assumed to be conservative in other taxa. In parallel, analogous variability in characters such as the shape and presence of spine I and dorsal sculpturing, has been also reported recently in Claxtonia spp.42.

On the other hand, genetic data, when taken in isolation, can also be deceiving, and this pertains specifically to the COI marker, often uncritically utilised and termed as a universal tardigrade barcode as (I) it is often



Figure 6. Habitus of *V. viridianus* (SEM): (**A**) dorsal view; (**B**) frontal view; (**C**) rear view. Scale bars = 50 µm.



Figure 7. Anterior body portion of *V. viridianus* (SEM): (A) dorsal view; (B) head. Scale bars = 20 µm.

difficult to amplify, and (II) may fail in delimiting species (Fig. 5;⁴³). Multiple markers are advisable⁴⁴, and both ITS markers are good predictors of intra- and interspecific differences, as they are congruent with a spectrum of morphological variability analysed on many animals. Altogether, this reinforces the necessity of the integrative approach in tardigrade studies¹⁷.

In the light of our findings, some generalisations can be made regarding the diversity and classification of *Viridiscus*:



Figure 8. Cuticle of *V. viridianus* (SEM): (A) epicuticular granules and micropores; (B) lateral portion of disrupted epicuticle; (C,D) close up of intracuticular pillars. Scale bars in µm.



Figure 9. Leg morphology of *V. viridianus* (PCM). Arrows indicate pedal platelets in central leg portions, white asterisks indicate distinctly demarcated, central oval areas in pedal platelets, and black asterisks indicate pulvini in proximal leg portions. Scale bars = $20 \mu m$.



Figure 10. Morphological details of *V. viridianus* (PCM): (A–C) conoid primary clava; (D) dactyloid primary clava; (E) typical for most echiniscids, tubby *Echiniscus*-like primary clava; (F) subcephalic plates; (G) claws I; (H) claws II; (I) claws III. Scale bars = 10 μ m.

- 1. The green body colour is a result of dorsal cuticular plate pigmentation, but orange and yellow (probably carotenoid-derived pigments), are present in all *Viridiscus* spp., as in most other echiniscids. Thus, two characteristics of the dorsal armour, the presence of the endocuticular sponge layer and green pigmentation, are advanced characters.
- 2. Although more variable than previously reported, the relative length of cirrus *A* with respect to the body length (*bo*) serves as a good criterion in morphological species delineation. The extreme elongation of cirrus



Figure 11. Leg structures of *V. viridianus* (SEM): (**A**) claws I; (**B**) claws II; (**C**) claws III; (**D**) claws IV; (**E**) leg morphology. Arrows indicate pedal platelets in central leg portions, white asterisks indicate distinctly demarcated, central oval areas in pedal platelets, and black asterisks indicate pulvini in proximal leg portions. Scale bars in µm.

| | | | | | | Analyses | | | | |
|-------------------------------|-------------|--------------------------------------|--|---|-----------------|-------------------------|-----|-----|--|--|
| Species | Sample code | Coordinates altitude | Locality | Sample type | Collector | РСМ | SEM | DNA | | |
| Viridiscus celatus sp. | US.078 | 36°18'N 82°22'W ca. 520 m asl | Tennessee, Washington County, Johnson City | Moss from a concrete cap on a brick fence post | Diane R. Nelson | 1a (♀) | - | 4a | | |
| nov.* | US.081 | 36°18'N 82°22'W ca. 520 m asl | Tennessee, Washington County, Johnson City | Moss from a concrete cap on a brick fence post | Diane R. Nelson | 6a (4♀+2♂), 4j | - | 4a | | |
| Viridiscus perviridis | US.086 | 33°12′51″N 87°34′17″W 43 m asl | Alabama, Tuscaloosa, campus of University of Alabama | Lichen** from rock boulder | Sogol Momeni | 7a (♀), 2j, 1l | - | 6a | | |
| | US.087 | 33°16'15"N 87°28'21"W 76 m asl | Alabama, vicinity of Lake Harris | Lichen*** from rock boulder | Sogol Momeni | 28a (16♀ + 12♂), 2v | - | 10a | | |
| | US.089 | 33°17′23″N 87°29′2″W 85 m asl | Alabama, vicinity of Lake Nicol | Lichen*** from rock boulder | Sogol Momeni | 23a (20♀ + 3♂), 11j, 11 | 40a | 10a | | |
| | US.090 | 33°16'11"N 87°28'12"W 67 m asl | Alabama, vicinity of Lake Harris | Lichen*** from rock boulder | Sogol Momeni | 3a (♀) | - | 4a | | |
| Viridiscus viridianus | US.092 | 33°17′23″N 87°29′1″W 85 m asl | Alabama, vicinity of Lake Nicol | Lichen*** from rock boulder | Sogol Momeni | 30a (27♀ + 3♂), 12j | 40a | 10a | | |
| | US.159 | 33°17'22"N 87°29'0"W 82 m asl | Alabama, vicinity of Lake Nicol | Lichen*** from rock | Sogol Momeni | 12a (10♀+2♂), 5j, 3l | 20a | 8a | | |
| | US.161 | 33°16'15"N 87°28'20"W 76 m asl | Alabama, vicinity of Lake Harris | Moss from rock | Sogol Momeni | 11a (10♀ + 1♂), 5j, 4l | 20a | 8a | | |
| | US.165 | 25°45'13"N 80°22'44"W 1 m asl | Florida, University Park | Lichen from tree | Jason Pienaar | 10a (6♀+4♂) | 10a | 8a | | |

Table 5. List of newly found populations used in analyses. Types of analyses: (PCM) imaging and morphometry in PCM, (SEM) imaging in SEM, (DNA) DNA sequencing. Number in each analysis indicates how many specimens were utilised in a given method (*a* adults, *v* exuvia, *j* juveniles, *l* larvae). **Viridiscus* aff. *viridianus* in ¹⁷. ***Flavoparmelia* sp. ****Flavoparmelia* baltimorensis.

A characterises V. perviridis (most often \gg 50% of bo), followed by cirri A longer than the echiniscid average (30–50% of bo) in V. celatus **sp. nov.** and V. viridissimus, and cirri A that can be classified as short (< 20% of bo) in V. clavispinosus **nom. inq.**, V. viridianus, and V. viridis²³.

- 3. Atypical morphs are probably present in populations of all *Viridiscus* spp. We recorded them in *V. perviridis*, *V. viridianus* (this study), and *V. viridissimus*¹⁷. This means that morphology alone does not offer a fully credible dataset for establishing new species.
- 4. The description of *V. viridis* by Pilato et al.²³ was based on only two individuals and likely does not reflect the spectrum of morphological variability of this species, but plainly indicates the specific nature of its dorsal sculpturing. The sculpturing embraces particularly widely spaced epicuticular granules, making it dissimilar to all other *Viridiscus* spp. Unless integratively redescribed from the Hawaiian Islands and later verified from other regions of the globe, it should be temporarily considered an endemic of this archipelago.

| | | Range | | Mean | | SD | | Holotype | |
|-------------------------------|---|------------|-------------|------|-------|------|------|----------|-------|
| Character | N | μm | sp | μm | sp | μm | sp | μm | sp |
| Body length | 5 | 195-252 | 449-533 | 218 | 502 | 21 | 32 | 252 | 516 |
| Scapular plate length | 5 | 38.0-48.8 | - | 43.5 | - | 4.2 | - | 48.8 | - |
| Head appendage lengths | | | | | | | | | |
| Cirrus internus | 5 | 9.6-11.8 | 23.2-27.4 | 10.7 | 24.7 | 0.9 | 2.0 | 11.4 | 23.4 |
| Cephalic papilla | 5 | 6.2-7.1 | 13.9–16.5 | 6.6 | 15.3 | 0.3 | 1.1 | 7.1 | 14.5 |
| Cirrus externus | 5 | 11.7-15.2 | 25.4-36.6 | 13.9 | 32.1 | 1.5 | 4.2 | 15.2 | 31.1 |
| Clava | 5 | 5.1-6.5 | 11.1–15.7 | 5.9 | 13.6 | 0.6 | 1.7 | 6.3 | 12.9 |
| Cirrus A | 5 | 70.5-105.0 | 152.9–243.6 | 86.6 | 199.8 | 15.1 | 32.9 | 100.1 | 205.1 |
| Cirrus A/body length ratio | 5 | 34-49% | - | 40% | - | 6% | - | 40% | - |
| Body appendage lengths | | | | | | | | | |
| Spine on leg I length | 5 | 2.1-3.7 | 4.9-8.7 | 3.0 | 6.8 | 0.7 | 1.5 | 3.7 | 7.6 |
| Papilla on leg IV length | 5 | 3.8-4.7 | 9.5-10.9 | 4.4 | 10.0 | 0.4 | 0.5 | 4.7 | 9.6 |
| Number of teeth on the collar | 5 | 6-10 | - | 8.2 | - | 1.5 | - | 10 | - |
| Claw I heights | | | | | | | | | |
| Branch | 4 | 14.3-17.2 | 33.0-37.6 | 15.4 | 35.4 | 1.3 | 1.9 | 17.2 | 35.2 |
| Spur | 3 | 1.5-1.8 | 3.7-4.4 | 1.7 | 4.0 | 0.2 | 0.3 | ? | ? |
| Spur/branch height ratio | 3 | 10-12% | - | 11% | - | 1% | - | ? | - |
| Claw II heights | • | | | | | | | | |
| Branch | 4 | 14.2-17.8 | 31.2-37.8 | 15.7 | 35.0 | 1.7 | 2.9 | 17.8 | 36.5 |
| Spur | 4 | 1.4-2.1 | 3.0-4.6 | 1.8 | 4.0 | 0.3 | 0.7 | 2.1 | 4.3 |
| Spur/branch height ratio | 4 | 10-13% | - | 11% | - | 2% | - | 12% | - |
| Claw III heights | | | | | | | | | |
| Branch | 5 | 12.3-17.0 | 29.7-37.8 | 14.6 | 33.6 | 2.0 | 3.0 | 17.0 | 34.8 |
| Spur | 3 | 1.6-1.8 | 3.5-3.9 | 1.7 | 3.7 | 0.1 | 0.2 | 1.8 | 3.7 |
| Spur/branch height ratio | 3 | 11-12% | - | 11% | - | 1% | - | 11% | - |
| Claw IV heights | | | | | | | | | |
| Branch | 5 | 15.3-21.4 | 34.9-45.9 | 17.7 | 40.8 | 2.7 | 4.6 | 21.4 | 43.9 |
| Spur | 4 | 1.6-2.3 | 3.9-4.7 | 1.9 | 4.4 | 0.3 | 0.4 | 2.3 | 4.7 |
| Spur/branch height ratio | 4 | 10-13% | - | 11% | - | 1% | - | 11% | - |

Table 6. Measurements (in μ m) of selected morphological structures of adult females of *Viridiscus celatus* **sp. nov.** mounted in Hoyer's medium. *sp* the proportion between the length of a given structure and the length of the scapular plate, ? unknown.

| | Allotype 🖒 | | Paratype 👌 | | Juvenile 1 | | Juvenile 2 | | Juvenile 3 | | Juvenile 4 | |
|-------------------------------|------------|------|------------|-------|------------|-------|------------|------|------------|-------|------------|------|
| Character | μm | sp | μm | sp | μm | sp | μm | sp | μm | sp | μm | sp |
| Body length | 208 | 470 | 189 | 471 | 187 | 475 | 154 | 524 | 185 | 532 | 152 | 466 |
| Scapular plate length | 44.3 | - | 40.1 | - | 39.4 | - | 29.4 | - | 34.8 | - | 32.6 | - |
| Head appendage lengths | | | | | | | | | | | | |
| Cirrus internus | 9.4 | 21.2 | 11.4 | 28.4 | 10.9 | 27.7 | 7.6 | 25.9 | 9.7 | 27.9 | 9.3 | 28.5 |
| Cephalic papilla | 6.3 | 14.2 | 5.5 | 13.7 | 5.9 | 15.0 | 5.1 | 17.3 | 5.5 | 15.8 | 4.5 | 13.8 |
| Cirrus externus | 10.9 | 24.6 | 15.4 | 38.4 | 12.3 | 31.2 | 8.2 | 27.9 | 11.7 | 33.6 | 11.8 | 36.2 |
| Clava | 5.2 | 11.7 | 5.0 | 12.5 | 5.9 | 15.0 | 4.3 | 14.6 | 5.7 | 16.4 | 4.8 | 14.7 |
| Cirrus A | ? | ? | 78.4 | 195.5 | 82.6 | 209.6 | ? | ? | 71.2 | 204.6 | ? | ? |
| Cirrus A/body length ratio | ? | - | 41% | - | 44% | - | ? | - | 38% | - | ? | - |
| Body appendage lengths | | | | | | | | | | | | |
| Spine on leg I length | 2.8 | 6.3 | 2.4 | 6.0 | 2.7 | 6.9 | 2.2 | 7.5 | 2.6 | 7.5 | 1.7 | 5.2 |
| Papilla on leg IV length | 4.5 | 10.2 | 3.7 | 9.2 | ; | ? | 3.3 | 11.2 | 3.8 | 10.9 | 3.3 | 10.1 |
| Number of teeth on the collar | 9.0 | - | 11.0 | - | ; | - | 7.0 | - | 9.0 | - | 8.0 | - |
| Claw I heights | | | | | | | | | | | | |
| Branch | 14.0 | 31.6 | 15.3 | 38.2 | 16.5 | 41.9 | 9.7 | 33.0 | 12.9 | 37.1 | 11.9 | 36.5 |
| Spur | 1.6 | 3.6 | 2.0 | 5.0 | 1.9 | 4.8 | 1.2 | 4.1 | 1.6 | 4.6 | 1.5 | 4.6 |
| Spur/branch height ratio | 11% | - | 13% | - | 12% | - | 12% | - | 12% | - | 13% | - |
| Claw II heights | | | | • | | | | | | | | |
| Branch | 13.2 | 29.8 | 14.6 | 36.4 | ? | ? | 10.0 | 34.0 | 12.3 | 35.3 | 10.9 | 33.4 |
| Spur | ? | ? | 1.7 | 4.2 | ? | ? | 1.3 | 4.4 | 1.7 | 4.9 | 1.4 | 4.3 |
| Spur/branch height ratio | ? | - | 12% | - | ? | - | 13% | - | 14% | - | 13% | - |
| Claw III heights | | | | | | | | | | | | |
| Branch | 13.4 | 30.2 | 14.7 | 36.7 | 14.3 | 36.3 | 9.6 | 32.7 | 12.1 | 34.8 | 10.7 | 32.8 |
| Spur | 1.5 | 3.4 | 1.9 | 4.7 | ? | ? | 1.3 | 4.4 | 1.6 | 4.6 | 1.4 | 4.3 |
| Spur/branch height ratio | 11% | - | 13% | - | ? | - | 14% | - | 13% | - | 13% | - |
| Claw IV heights | | | | | | | | | | | | |
| Branch | 15.8 | 35.7 | 15.9 | 39.7 | 17.9 | 45.4 | 11.6 | 39.5 | 14.5 | 41.7 | 14.6 | 44.8 |
| Spur | 2.0 | 4.5 | ? | ? | ? | ? | ? | ? | 1.8 | 5.2 | ? | ? |
| Spur/branch height ratio | 13% | - | ? | - | ? | - | ? | - | 12% | - | ? | - |

Table 7. Measurements (in μ m) of selected morphological structures of adult males and juveniles of *Viridiscus celatus* **sp. nov.** mounted in Hoyer's medium. *sp* the proportion between the length of a given structure and the length of the scapular plate, ? unknown.



Figure 12. Holotype of V. celatus sp. nov. (PCM, female, dorsal view). Scale bar = $20 \mu m$.



Figure 13. Type specimens of *V. celatus* **sp. nov.** (PCM): (**A**) allotype (male, dorsal view); (**B**) paratype (female, dorsolateral view). Arrowheads indicate rudimentary papillae on legs II–III. Scale bars = 20 μm.



Figure 14. Morphological details of *V. celatus* **sp. nov.** (PCM): (**A**) subcephalic plates (holotype); (**B**) claws III (paratype, juvenile); (**C**) claws IV (holotype). Scale bars = 10 µm.

Conclusions

The extent of intraspecific morphological variability within *Viridiscus* is considerable and exceeds the usual variation encountered within echiniscid populations. Atypical morphs of *Viridiscus* species are not linked with sexual dimorphism, phenology, or ontogeny, and seem to represent idiopathic deviations from the most common morphs. *Viridiscus celatus* **sp. nov.**, found only in association with large populations of *V. viridissimus*, is described integratively based on genetics and morphology. In addition, the validity of *V. clavispinosus* is questioned, and its synonymy with *V. viridianus* is implied. Males were reported in populations of *Viridiscus celatus* **sp. nov.**, *V. viridianus*, and *V. viridissimus* so far.



Figure 15. Habitats harbouring *Viridiscus* populations in *Andreaea* mosses growing on boulders by Lake Harris, Tuscaloosa, Alabama: (**A**) an overall view of the rock substrate; (**B**) close up of the moss growing on the vertical surfaces of the boulder (arrow indicates the moss matt); (**C**) close up of the pleurocarpous moss and leaf structure when dry; (**D**) moss leaves seen under a stereomicroscope (arrows indicate animals on moss leaves in the tun state).

| Species | Population | 18S rRNA | 28S rRNA | ITS-1 | ITS-2 | COI |
|-----------------------------|------------|----------------|---------------|----------------------|--------------------|------------------|
| Viridicaus calatus on nou* | US.078 | MZ868197 | OK094230 | OK094211-4 | OK094173-6 | MZ852064-6 |
| viriaiscus ceiaius sp. nov. | US.081 | MZ868198 | OK094231 | OK094215-8 | OK094177-80 | MZ852067-9 |
| | PT.042 | MK529696 | MK529726-7 | OK094219-20 | OK094181-2 | - |
| Viridiscus perviridis | US.086 | OR519990-2 | OR520001-3 | - | OR520055-7 | OR502563-5 |
| | VN.028 | MZ868199 | OK094232 | OK094221-3 | OK094183-5 | - |
| | US.087 | OR519993-4 | OR520004-5 | OR520012-8 | OR520058-64 | OR502566-72 |
| | US.089 | - | - | OR520019-27 | OR520065-73 | OR502573-81 |
| | US.090 | - | - | OR520028 | OR520074 | OR502582 |
| Viridiscus viridianus | US.092 | - | - | OR520029-32 | OR520075-8 | OR502583-6 |
| | US.159 | OR519995-6 | OR520006-7 | OR520033-9 | OR520079-85 | OR502587-93 |
| | US.161 | OR519997-8 | OR520008-9 | OR520040-7 | OR520086-93 | OR502594-601 |
| | US.165 | OR519999-20000 | OR520010-1 | OR520048-54 | OR520094-100 | OR502602-8 |
| | US.078 | MZ868191-3 | OK094224, 6–7 | OK094186-90, 207-208 | OK094148-52, 69-70 | MZ852046-9, 62-3 |
| Viridiscus viridissimus | US.081 | MZ868194 | OK094225 | OK094191-206 | OK094153-68 | MZ852050-61 |
| | VN.028 | MZ868195-6 | OK094228-9 | OK094209-10 | OK094171-2 | - |
| Echiniscus succineus | MG.005 | MK675903 | MK675914 | MT374198 | MK675925 | MK649675 |

Table 8. GenBank accession numbers for the sequences analysed in this work. New sequences in bold.

 *Viridiscus aff. viridianus in¹⁷.

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Materials and methods Sample collection and processing

Previous sampling experience^{31,45} showed that *Viridiscus* species are often found in mosses and lichens growing on rapidly draining vertical surfaces (see e.g.⁴⁶) such as the sides of rocks (Fig. 15), gravestones, and tree trunks in human-altered environment. Therefore, moss and lichen samples were collected from several localities in Alabama, Florida, and Tennessee from February 2020 to May 2022 (Table 5). All samples were stored in acid-free paper envelopes and allowed to dry overnight in an air-conditioned room at 25 °C and 50% relative humidity. The air-dried samples were suspended over deionised water using the custom-built Baermann pan described by Davison⁴⁷ for at least 12 h to separate the motile meiofauna from the moss and lichen particles. The deionised water was then filtered through 25 µm mesh and the sieve was rinsed into 30 mm Petri dishes with locally collected rainwater to search for tardigrades. The air-dried lichens were soaked in deionised water for at least 12 h, active tardigrades were extracted with an Irwin loop and transferred into 30 mm Petri dishes containing rainwater.

Microscopy and imaging

Individual tardigrades from the samples were visually inspected for diagnostic traits using an Olympus BX53 phase contrast microscope (PCM) associated with an Olympus DP74 digital camera. All specimens were divided into morphological and molecular analyses (details in Table 5). The excess of individuals was frozen to store for future analyses. For morphology and morphometry, individuals were permanently mounted on microscope slides in Hoyer's medium. Dried slides were sealed with nail polish and examined under the Olympus BX53 PCM. Specimens for imaging in the SEM were CO_2 critical point-dried, coated with gold and examined in the Versa 3D DualBeam SEM at the ATOMIN facility of the Jagiellonian University. All figures were assembled in Corel Photo-Paint X8. For deep structures that could not be fully focused on a single PCM photograph, a series of images was taken every *ca.* 0.1 mm of vertical focusing and then assembled manually in Corel Photo-Paint into a single deep-focus image.

Morphometry

All individuals of the new species from Tennessee and selected individuals of *V. viridianus* from Alabama were chosen for morphometry. The measurements are in micrometres (μ m), and only undamaged structures with the appropriate orientation were used. The body length measurement is from the anterior to the posterior ends of the body, excluding the hind limbs. The *bo*% was calculated for cirrus *A*, which is the ratio of cirrus *A* length to the body, and *sp*% is the ratio of the length of the given structure to the length of the scapular plate⁴⁸. Morphometric data were handled using the Echiniscoidea ver. 1.4 template available from the Tardigrada Register, http://www.tardigrada.net/register ⁴⁹.

Comparative material

The morphology of *Viridiscus* spp. was compared with paratypes of *V. viridianus* and syntypes of *V. perviridis* (deposited in the Civic Museum of Natural History in Verona and University of Modena and Reggio Emilia, Italy), and original descriptions or redescriptions of species^{17,23-26}. Slides of *V. viridissimus* from Tennessee, including the "*miraviridis* morphotype" (¹⁶, synonymised with *V. viridissimus* by Gąsiorek et al.¹⁷), were also examined.

Genotyping

A Chelex 100 resin (Bio-Rad) extraction method was used for DNA extraction^{50,51}. Hologenophores were recovered after the extraction and mounted on permanent slides in Hoyer's medium in most cases; in other cases, paragenophores were preserved⁵²; all are deposited in the Faculty of Biology, Jagiellonian University in Kraków. Five DNA fragments (nuclear markers: 18S rRNA, 28S rRNA, ITS-1, ITS-2; mitochondrial marker: COI) were amplified and sequenced according to the protocols described in⁵¹; primers and original references for specific PCR programmes are listed in Supplementary Material 3. GenBank accession numbers for the species used in calculating phylogenies and the genetic framework (mainly based on the dataset from Gasiorek et al.¹⁷) are provided in Table 8. ITS and COI sequences were separately aligned with sequences from Echiniscus succineus Gasiorek & Vončina, 2019⁵³ as an outgroup using the ClustalW Multiple Alignment tool⁵⁴ implemented in BioEdit ver. 7.2.5⁵⁵. The remaining gaps were left intact in ITS alignments. The 18S and 28S rRNA gene fragments were not used for the species delimitation purposes but are provided for future broader-scale phylogenies.

Phylogeny

The sequences of the ITS fragments were concatenated to generate a matrix of 1058 bp in SequenceMatrix⁵⁶. Using PartitionFinder version 2.1.157 with applied Bayesian Information Criterion (BIC) and the greedy algorithm⁵⁸, the best substitution model (the lowest BIC) and partitioning scheme were chosen for posterior phylogenetic analysis. As the best-fit partitioning scheme, PartitionFinder indicated one partition with the best-fitted model GTR+G. Bayesian inference (BI) marginal posterior probabilities were calculated using MrBayes v.3.2⁵⁹. Random starting trees were used, and the analysis was run for ten million generations, sampling the Markov chain every 1000 generations. An average standard deviation of split frequencies of < 0.01 was used as a guide to ensure that the two independent analyses had converged. Tracer v1.360 was then used to ensure Markov chains had reached stationarity and to determine the correct burn-in' for the analysis, in this case the first 10% of generations. The Effective Sample Size values were greater than 200, and the consensus tree was obtained after summarising the resulting topologies and discarding the 'burn-in'.

ModelFinder⁶¹ was used to choose the best-fit models for two partitions in Maximum Likelihood (ML): TIM3e+G4 (ITS-1) and K2P+G4 (ITS-2), chosen according to the BIC. W-IQ-TREE was used for ML reconstruction^{62,63}. One thousand ultrafast bootstrap (UFBoot) replicates were applied to provide support values for branches⁶⁴.

Both ITS and COI alignments were uploaded separately to the Assemble Species by Automatic Partitioning (ASAP) webpage⁶⁵, and automatic barcode gap discovery (ABGD) web⁶⁶ to obtain six independent marker-based primary species hypotheses using uncorrected pairwise distances. The partitions with the lowest ASAP and ABGD scores and *p* values < 0.05 were chosen as the best-fit hypotheses. Finally, Bayesian Poisson tree processes (bPTP⁶⁷) were applied to the ML phylogenetic trees run on three markers separately in W-IQ-TREE (COI bestfitted model: HKY + F + G4). The calculations were conducted with 100,000 MCMC generations, thinning the set to 100, with 10% burn-in, and with searches for maximum likelihood and Bayesian solutions. All final consensus trees were visualised by FigTree v.1.4.3 available from https://tree.bio.ed.ac.uk/software/figtree.

Data availability

Data generated or analysed during this study are included in the published article and its supplementary information files. All sequences are deposited in GenBank. The publication was registered in ZooBank: urn:lsid:zoobank. org:pub:E2AA1924-C159-460B-9C8B-F23D951A5FA3.

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References

- 1. Wu, R., Pisani, D. & Donoghue, P. C. J. The unbearable uncertainty of panarthropod relationships. Biol. Lett. 19(1), 20220497. https://doi.org/10.1098/rsbl.2022.0497 (2023).
- 2. Degma, P. & Guidetti, R. Actual checklist of Tardigrada species, 42nd edn. Accessed 15th April 2023. https://doi.org/10.25431/ 11380_1178608 (2023).
- 3. Marcus, E. Zur Anatomie und Ökologie mariner Tardigraden. Zoologische Jahrbücher abteilung für Systematik Ökologie und Geographie der Tiere 53, 487-558 (1927).
- 4. Kristensen, R. M. Generic revision of the Echiniscidae (Heterotardigrada), with a discussion of the origin of the family. In Biology of Tardigrades. Selected Symposia and Monographs U.Z.I Vol. 1 (ed. Bertolani, R.) 261-335 (Mucchi, 1987).
- 5. Fontoura, P., Bartels, P. J., Jørgensen, A., Kristensen, R. M. & Hansen, J. G. A dichotomous key to the genera of the marine heterotardigrades (Tardigrada). *Zootaxa* **4294**(1), 1–45. https://doi.org/10.11646/zootaxa.4294.1.1 (2017). 6. Bertolani, R. *et al.* Phylogeny of Eutardigrada: New molecular data and their morphological support lead to the identification of
- new evolutionary lineages. Mol. Phylogenet. Evol. 76, 110-126. https://doi.org/10.1016/j.ympev.2014.03.006 (2014).
- 7. Nelson, D. R., Guidetti, R. & Rebecchi, L. Phylum Tardigrada, Chapter 17. In Ecology and General Biology: Thorp and Covich's Freshwater Invertebrates (eds Thorp, J. & Rogers, D. C.) 347-380 (Academic Press, 2015). https://doi.org/10.1016/B978-0-12-385026-3.00017-6.
- 8. Nelson, D. R., Bartels, P. J. & Guil, N. Tardigrade ecology, Chapter 7. In Water Bears: The Biology of Tardigrades. Zoological Monographs (ed. Schill, R. O.) 163-210 (Springer, 2018). https://doi.org/10.1007/978-3-319-95702-9_7.
- 9. Jørgensen, A. Graphical presentation of the African tardigrade fauna using GIS with the description of Isohypsibius malawiensis sp. n. (Eutardigrada: Hypsibiidae) from Lake Malawi. Zool. Anzeiger 240(3-4), 441-450. https://doi.org/10.1078/0044-5231-00052 (2001).
- 10. Claxton, S. K. The taxonomy and distribution of Australian terrestrial tardigrades, PhD dissertation (Macquarie University, 2004). 11. Gąsiorek, P. et al. Echiniscus virginicus complex: The first case of pseudocryptic allopatry and pantropical distribution in tardigrades.
- Biol. J. Linn. Soc. 128(4), 789-805. https://doi.org/10.1093/biolinnean/blz147 (2019)
- 12. Meyer, H. A. Terrestrial and freshwater Tardigrada of the Americas. Zootaxa 3747(1), 1-71. https://doi.org/10.11646/zootaxa. 3747.1.1 (2013).

- Kaczmarek, Ł, Michalczyk, Ł & McInnes, S. J. Annotated zoogeography of non-marine Tardigrada. Part III: North America and Greenland. Zootaxa 4203(1), 1–249. https://doi.org/10.11646/zootaxa.4203.1.1 (2016).
- Degma, P. & Guidetti, R. Tardigrade taxa, Chapter 15. In Water Bears: The Biology of Tardigrades. Zoological Monographs (ed. Schill, R. O.) 371-409 (Springer, 2018).
- Gąsiorek, P., Morek, W., Stec, D. & Michalczyk, Ł. Untangling the *Echiniscus* Gordian knot: Paraphyly of the "arctomys group" (Heterotardigrada: Echiniscidae). Cladistics 35(6), 633–653. https://doi.org/10.1111/cla.12377 (2019).
- Nelson, D. R. et al. Two new species of Tardigrada from moss cushions (Grimmia sp.) in a xerothermic habitat in northeast Tennessee (USA, North America), with the first identification of males in the genus Viridiscus. PeerJ 8, e10251. https://doi.org/10.7717/peerj.10251 (2020).
- Gasiorek, P., Vončina, K., Nelson, D. R. & Michalczyk, Ł. The importance of being integrative: A remarkable case of synonymy in the genus *Viridiscus* (Heterotardigrada: Echiniscidae). *Zool. Lett.* 7, 13. https://doi.org/10.1186/s40851-021-00181-z (2021).
- Ramazzotti, G. Il gruppo dell'Echiniscus viridis con la nuova specie E. perviridis e Macrobiotus pustulatus altra nuova specie (Tardigrada). Atti della Società Italiana di Scienze Naturali e del Museo Civico di Storia Naturale in Milano 98(4), 303–309 (1959).
- 19. Murray, J. Tardigrada. British Antarctic Expedition 1907–1909. Reports on the Scientific Investigations, Biology (Part V) 1, 83–187 (1910).
- 20. Péterfi, F. Contribuțiuni la cunoașterea tardigradelor din R. P. R. Studii și cercetări de Biologie 7, 149-155 (1956).
- 21. du Bois-Reymond Marcus, E. Sobre tardígrados brasileiros. *Comunicaciones zoológicas del Museo de Historia Natural de Montevideo* 1(13), 1–19 (1944).
- 22. Bonifacio, A., Guidetti, R., Altiero, T., Sergo, V. & Rebecchi, L. Nature, source and function of pigments in tardigrades: *In vivo* Raman imaging of carotenoids in *Echiniscus blumi*. *PLoS One* 7, 11. https://doi.org/10.1371/journal.pone.0050162 (2012).
- Pilato, G., Fontoura, P. & Lisi, O. Remarks on the *Echiniscus viridis* group, with the description of a new species (Tardigrada, Echiniscidae). J. Limnol. 66(Suppl. 1), 33–39. https://doi.org/10.4081/jlimnol.2007.s1.33 (2007).
- Fontoura, P., Pilato, G. & Lisi, O. Tardigrada from Santo Antão Island (Archipelago of Cape Verde, West Africa) with the description of a new species. Zootaxa 2838(1), 30–40. https://doi.org/10.11646/zootaxa.2838.1.2 (2011).
- Rocha, A. *et al.* Actual state of knowledge of the limno-terrestrial tardigrade fauna of the Republic of Argentina and new genus assignment for *Viridiscus rufoviridis* (du Bois-Reymond Marcus, 1944). *Diversity* 15, 222. https://doi.org/10.3390/d15020222 (2023).
- Pilato, G., Fontoura, P. & Lisi, O. New description of *Echiniscus viridis* Murray, 1910 and remarks on the viridis group. New Zeal. J. Zool. 35(1), 85–92. https://doi.org/10.1080/03014220809510105 (2008).
- Nelson, D. R. & Adkins, R. G. Distribution of tardigrades within a moss cushion: Do tardigrades migrate in response to changing moisture conditions? *Zool. Anz.* 240(3–4), 493–500. https://doi.org/10.1078/0044-5231-00058 (2001).
- Miller, W. R. *et al.* Tardigrades of North America: New records of occurrence for three species of green tardigrades (Heterotardigrada, Echiniscoidea, Echiniscidae, *Viridiscus*). *Trans. Kans. Acad. Sci.* 124(1–2), 235–241. https://doi.org/10.1660/062.123.0122 (2020).
- 29. Fontoura, A. P. Primeiro contributo para o conhecimento da fauna de tardigrados dos Açores. Arquipélago. Série Ciências da Natureza 6, 45–56 (1985).
- Fontoura, P., Pilato, G. & Lisi, O. Echiniscidae (Tardigrada, Heterotardigrada) from Faial and Pico Islands, the Azores, with the description of two new species. Zootaxa 1693(1), 49–61. https://doi.org/10.11646/zootaxa.1693.1.4 (2008).
- Momeni, S., Fuentes-González, J. & Pienaar, J. How to culture limnoterrestrial heterotardigrades. *Invertebr. Biol.* 141(1), e12360. https://doi.org/10.1111/ivb.12360 (2022).
- Maucci, W. A contribution to the knowledge of North American Tardigrada with emphasis on the fauna of Yellowstone National Park (Wyoming). In *Biology of Tardigrades. Selected Symposia and Monographs U.Z.I* Vol. 1 (ed. Bertolani, R.) 187–210 (Mucchi, 1987).
- Grigarick, A. A., Schuster, R. O. & Nelson, D. R. Heterotardigrada of Venezuela (Tardigrada). Pan Pac. Entomol. 59(1-4), 64-77 (1983).
- Jörger, K. M., Norenburg, J. L., Wilson, N. G. & Schrödl, M. Barcoding against a paradox? Combined molecular species delineations reveal multiple cryptic lineages in elusive meiofaunal sea slugs. *BMC Evol. Biol.* 12, 245. https://doi.org/10.1186/1471-2148-12-245 (2012).
- Rossberg, A. G., Rogers, T. & McKane, A. J. Are there species smaller than 1 mm?. Proc. R. Soc. B 280(1767), 20131248. https:// doi.org/10.1098/rspb.2013.1248 (2013).
- Mąkol, J., Saboori, A. & Felska, M. Inter- and intraspecific variability of morphological and molecular characters in Allothrombium species, with special reference to Allothrombium fuliginosum. Exp. Appl. Acarol. 78, 485–504. https://doi.org/10.1007/s10493-019-00400-5 (2019).
- Karanovic, T. Cladistic and quantitative shape analyses of five new syntopic Sarsamphiascus (Copepoda, Harpacticoida): Problems and solutions for diosaccin systematics and taxonomy. Syst. Biodivers. 18(8), 810–833. https://doi.org/10.1080/14772000.2020. 1832605 (2020).
- Morek, W., Surmacz, B., López-López, A. & Michalczyk, Ł. "Everything is not everywhere": Time-calibrated phylogeography of the genus Milnesium (Tardigrada). Mol. Ecol. 30(14), 3590–3609. https://doi.org/10.1111/mec.15951 (2021).
- Gąsiorek, P., Blagden, B. & Michalczyk, Ł. Towards a better understanding of echiniscid intraspecific variability: A redescription of *Nebularmis reticulatus* (Murray, 1905) (Heterotardigrada: Echiniscoidea). *Zool. Anz.* 283, 242–255. https://doi.org/10.1016/j. jcz.2019.08.003 (2019).
- 40. Gąsiorek, P. et al. Echiniscidae (Heterotardigrada) of South Africa. Zootaxa 5156(1), 1–238. https://doi.org/10.11646/zootaxa. 5156.1.1 (2022).
- Jørgensen, A., Møbjerg, N. & Kristensen, R. M. Phylogeny and evolution of the Echiniscidae (Echiniscoidea, Tardigrada)—An investigation of the congruence between molecules and morphology. J. Zool. Syst. Evol. Res. 49(1), 6–16. https://doi.org/10.1111/j. 1439-0469.2010.00592.x (2011).
- Gąsiorek, P., Degma, P. & Michalczyk, Ł. Hiding in the Arctic and in mountains: A (dis)entangled classification of *Claxtonia* (Heterotardigrada: Echiniscidae). *Zool. J. Linn. Soc.* https://doi.org/10.1093/zoolinnean/zlad029 (2023).
- Ranasinghe, U. G. S. L. *et al.* Multiple species delimitation approaches with COI barcodes poorly fit each other and morphospecies—An integrative taxonomy case of Sri Lankan Sericini chafers (Coleoptera: Scarabaeidae). *Ecol. Evol.* 12(5), e8942. https:// doi.org/10.1002/ece3.8942 (2022).
- 44. Zhang, H., Ning, X., Yu, X. & Bu, W.-J. Integrative species delimitation based on COI, ITS, and morphological evidence illustrates a unique evolutionary history of the genus *Paracercion* (Odonata: Coenagrionidae). *PeerJ* 9, e11459. https://doi.org/10.7717/peerj. 11459 (2021).
- Dewel, R. A., Dewel, W. C. & Roush, B. G. Unusual cuticle-associated organs in the heterotardigrade, *Echiniscus viridissimus. J. Morphol.* 212(2), 123–140. https://doi.org/10.1002/jmor.1052120204 (1992).
- Turberville, C. M., Fuentes-González, J. A., Rogers, S. & Pienaar, J. Moss phyllid morphology varies systematically with substrate slope. *Plant Ecol. Evol.* 154(3), 419–431. https://doi.org/10.5091/plecevo.2021.1839 (2021).
- Davison, P. G. Baermann pans. https://tardigradehunters.weebly.com/uploads/2/5/9/3/25930056/baermann_pan_instructions201 4breif.pdf (The International Society of Tardigrade Hunters, 2015).

- Dastych, H. Eine Neue Art der Gattung Mopsechiniscus Du Bois-Reymond Marcus, 1944 (Tardigrada) aus den Anden von Venezuela. Acta Biol. Benrodis 10, 91–101 (1999).
- Michalczyk, Ł & Kaczmarek, Ł. The Tardigrada Register: A comprehensive online data repository for tardigrade taxonomy. J. Limnol. 72(Suppl. 1), 175–181. https://doi.org/10.4081/jlimnol.2013.s1.e22 (2013).
- Casquet, J. T., Thebaud, C. & Gillespie, R. G. Chelex without boiling, a rapid and easy technique to obtain stable amplifiable DNA from small amounts of ethanol-stored spiders. *Mol. Ecol. Resour.* 12(1), 136–141. https://doi.org/10.1111/j.1755-0998.2011.03073.x (2012).
- Stec, D., Kristensen, R. M. & Michalczyk, Ł. An integrative description of *Minibiotus ioculator* sp. nov. from the Republic of South Africa with notes on *Minibiotus pentannulatus* Londoño et al., 2017 (Tardigrada: Macrobiotidae). Zool. Anz. 286, 117–134. https:// doi.org/10.1016/j.jcz.2020.03.007 (2020).
- Pleijel, F. et al. Phylogenies without roots? A plea for the use of vouchers in molecular phylogenetic studies. *Mol. Phylogenet. Evol.* 48(1), 369–371. https://doi.org/10.1016/j.ympev.2008.03.024 (2008).
- Gąsiorek, P. & Vončina, K. New Echiniscidae (Heterotardigrada) from Amber Mountain (Northern Madagascar). Evol. Syst. 3, 29–39. https://doi.org/10.3897/evolsyst.3.33580 (2019).
- Thompson, J. D., Higgins, D. G. & Gibson, T. J. CLUSTAL W: Improving the sensitivity of progressive multiple sequence alignment through sequence weighting, position-specific gap penalties and weight matrix choice. *Nucleic Acids Res.* 22(22), 4673–4680. https://doi.org/10.1093/nar/22.22.4673 (1994).
- Hall, T. A. BioEdit: A user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT. Nucleic Acids Symp. Ser. 41, 95–98 (1999).
- Vaidya, G., Lohman, D. J. & Meier, R. SequenceMatrix: Concatenation software for the fast assembly of multi-gene datasets with character set and codon information. *Cladistics* 27(2), 171–180. https://doi.org/10.1111/j.1096-0031.2010.00329.x (2011).
- Lanfear, R., Frandsen, P. B., Wright, A. M., Senfeld, T. & Calcott, B. PartitionFinder 2: New methods for selecting partitioned models of evolution for molecular and morphological phylogenetic analyses. *Mol. Biol. Evol.* 34(3), 772–773. https://doi.org/10. 1093/molbev/msw260 (2017).
- Lanfear, R., Calcott, B., Ho, S. Y. & Guindon, S. PartitionFinder: Combined selection of partitioning schemes and substitution models for phylogenetic analyses. *Mol. Biol. Evol.* 29(6), 1695–1701. https://doi.org/10.1093/molbev/mss020 (2012).
- Ronquist, F. & Huelsenbeck, J. P. MrBayes 3: Bayesian phylogenetic inference under mixed models. *Bioinformatics* 19(12), 1572– 1574. https://doi.org/10.1093/bioinformatics/btg180 (2003).
- 60. Rambaut, A., Suchard, M. A., Xie, D. & Drummond, A. J. Tracer v1.6. https://beast.bio.ed.ac.uk/Tracer (2014).
- 61. Kalyaanamoorthy, S., Minh, B. Q., Wong, T. K. F., von Haeseler, A. & Jermiin, L. S. ModelFinder: Fast model selection for accurate
- phylogenetic estimates. *Nat. Methods* 14, 587–589. https://doi.org/10.1038/nmeth.4285 (2017).
 62. Nguyen, L.-T., Schmidt, H. A., von Haeseler, A. & Minh, B. Q. IQ-TREE: A fast and effective stochastic algorithm for estimating maximum likelihood phylogenies. *Mol. Biol. Evol.* 32(1), 268–274. https://doi.org/10.1093/molbev/msu300 (2015).
- Trifinopoulos, J., Nguyen, L.-T., von Haeseler, A. & Minh, B. Q. W-IQ-TREE: A fast online phylogenetic tool for maximum likelihood analysis. *Nucleic Acids Res.* 44(W1), 232–235. https://doi.org/10.1093/nar/gkw256 (2016).
- Hoang, D. T., Chernomor, O., von Haeseler, A., Minh, B. Q. & Vinh, L. S. UFBoot2: Improving the ultrafast bootstrap approximation. Mol. Biol. Evol. 35(2), 518–522. https://doi.org/10.1093/molbev/msx281 (2018).
- Puillandre, N., Brouillet, S. & Achaz, G. ASAP: Assemble species by automatic partitioning. Mol. Ecol. Resour. 21(2), 609–620. https://doi.org/10.1111/1755-0998.13281 (2021).
- Puillandre, N., Lambert, A., Brouillet, S. & Achaz, G. ABGD, automatic barcode gap discovery for primary species delimitation. *Mol. Ecol.* 21(8), 1864–1877. https://doi.org/10.1111/j.1365-294X.2011.05239.x (2012).
- Zhang, J., Kapli, P., Pavlidis, P. & Stamatakis, A. A general species delimitation method with applications to phylogenetic placements. *Bioinformatics* 29(22), 2869–2876. https://doi.org/10.1093/bioinformatics/btt499 (2013).

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Author contributions

S.M. established the cultures of *V. viridianus*, collected molecular data, and wrote the first draft of the manuscript; P.G. collected molecular, morphometric and morphological data, analysed data, prepared figures, and wrote the manuscript; J.L. collected populations of *V. viridianus* in the field; S.C. collected molecular data; D.R.N. and R.A.F. collected populations of *V. celatus* **sp. nov.** and wrote the manuscript; J.P. wrote the manuscript. All authors read and approved the final version of the manuscript.

Competing interests

The authors declare no competing interests.

Additional information

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