

SHORT COMMUNICATION

TWO INVALID GENERA IN THE FAMILY BYTHINELLIDAE LOCARD, 1893 (CAENOGASTROPODA: TRUNCATELLOIDEA)

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ABSTRACT: Within the genus *Bythinella* Moquin-Tandon, 1856, with about 250 nominal species, the separation of several genera would be helpful in understanding the phylogenetic relationships between these snails, but neither morphology nor molecular loci studied so far can be used to distinguish any natural taxon over the range of species within this genus. Apart from *Bythinella* there are two other genera listed in the Bythinellidae by WoRMS: *Terrestrabythinella* Sitnikova, Starobogatov et Anistratenko, 1992, and *Strandzhia* Georgiev et Glöer, 2013. Considering the cytochrome c oxidase subunit I (COI) sequences, as well as poor morphological data, *Terrestrabythinella* should be considered a junior subjective synonym of *Bythinella*. In the case of *Strandzhia*, both COI and 18S sequences studied by us, as well as morphology re-examined (misinterpretation of the penis morphology in the original description) clearly show the assignment of *Strandzhia* to the genus *Grossuana* Radoman, 1983, thus *Strandzhia* becomes a junior objective synonym of *Grossuana*.

KEY WORDS: *Bythinella*; *Terrestrabythinella*; *Strandzhia*; morphology; mtDNA; synonyms

Bythinella Moquin-Tandon, 1856, with its type species *Bulimus viridis* Poiret, 1801, belongs to the family Bythinellidae Locard, 1893. For a long time, it was classified within the Amnicolidae Tryon, 1863, but both morphology (SZAROWSKA 2006) and molecular data (WILKE et al. 2001, 2013) confirmed the distinctness of the Bythinellidae. There are about 250 nominal species of *Bythinella* (WORMS 2024). However, neither morphology (GIUSTI & PEZZOLI 1977, FALNIOWSKI 1987, 2018, MAZAN 2000, HAASE et al. 2007, JASZCZYŃSKA in press), nor molecular loci studied so far (BICHAIN et al. 2007a, b, BENKE et al. 2009, 2011, FALNIOWSKI et al. 2009a, WILKE et al. 2010, FALNIOWSKI & SZAROWSKA 2011, SZAROWSKA

et al. 2016, JASZCZYŃSKA in press) may be used to separate any natural taxon of the genus level within the *Bythinella*.

SITNIKOVA et al. (1992) described from Ukraine a genus *Terrestrabythinella* Sitnikova, Starobogatov et Anistratenko, 1992, with its type species *T. baidashnikovi* Sitnikova, Starobogatov et Anistratenko, 1992. The other species: *T. carpathica* Sitnikova, Starobogatov et Anistratenko, 1992 was described and new family Terrestrabythinellidae Sitnikova, Starobogatov et Anistratenko, 1992 was created in the same publication (SITNIKOVA et al. 1992). Later, the third species of *Terrestrabythinella*: *T. amphibiotica* Anistratenko, 1995 was described (ANISTRATENKO 1995).



According to SITNIKOVA et al. (1992) there is no tubular penial gland in the male reproductive organs of *Terrestrabythinella*. There should be said that in such case the presence of a big flagellum would be enigmatic, and most probably the tubular gland was small and thus overlooked. In the female reproductive organs, according to SITNIKOVA et al. (1992), there was a diauly [meaning the existence of a distinct sperm duct (DAVIS 1967, HERSLER & PONDER 1998) – resulting in two separate female genital openings]. The diauly is characteristic for the Amnicolidae, but not for the Bythinellidae, since in the latter very broad folds forming the ventral channel mimic a spermathecal duct (SZAROWSKA 2006). According to SITNIKOVA et al. (1992), there is also a somewhat curious bursa with duct, and no receptaculum seminis. In fact, both bursa and receptaculum are extremely variable in *Bythinella* (e.g. GIUSTI & PEZZOLI 1977, FALNIOWSKI 1987, 2018, MAZAN 2000), the receptaculum may be as small as to be easily overlooked.

Considering the poor fidelity of the reproduction of the drawings of the shells and reproductive organs of *Falniowskia neglectissima* (Falniowski et Šteffek, 1989), the type species of *Falniowskia* Bernasconi, 1990, re-drawn from the original description (FALNIOWSKI & ŠTEFFEK 1989), the drawings and descriptions of SITNIKOVA et al. (1992) are thus even less convincing. Thus, these morphological characters alone may not constitute the base of systematics in the case of *Bythinella* and other truncatelloidean gastropods (FALNIOWSKI 2018). Molecular data are therefore necessary for clearer taxonomic conclusions.

For *Terrestrabythinella* we performed molecular data analysis following standard methods (FALNIOWSKI et al. 2023). The cytochrome c oxidase subunit I (COI) – a marker commonly used in animals' barcoding – definitely placed *Terrestrabythinella* within the *Bythinella*, as two species (mOTU A and B; Fig. 1), forming a sister clade with *B. viseuiana* Falniowski, Szarowska et Sirbu, 2009 (FALNIOWSKI et al. 2009b; mOTU C),

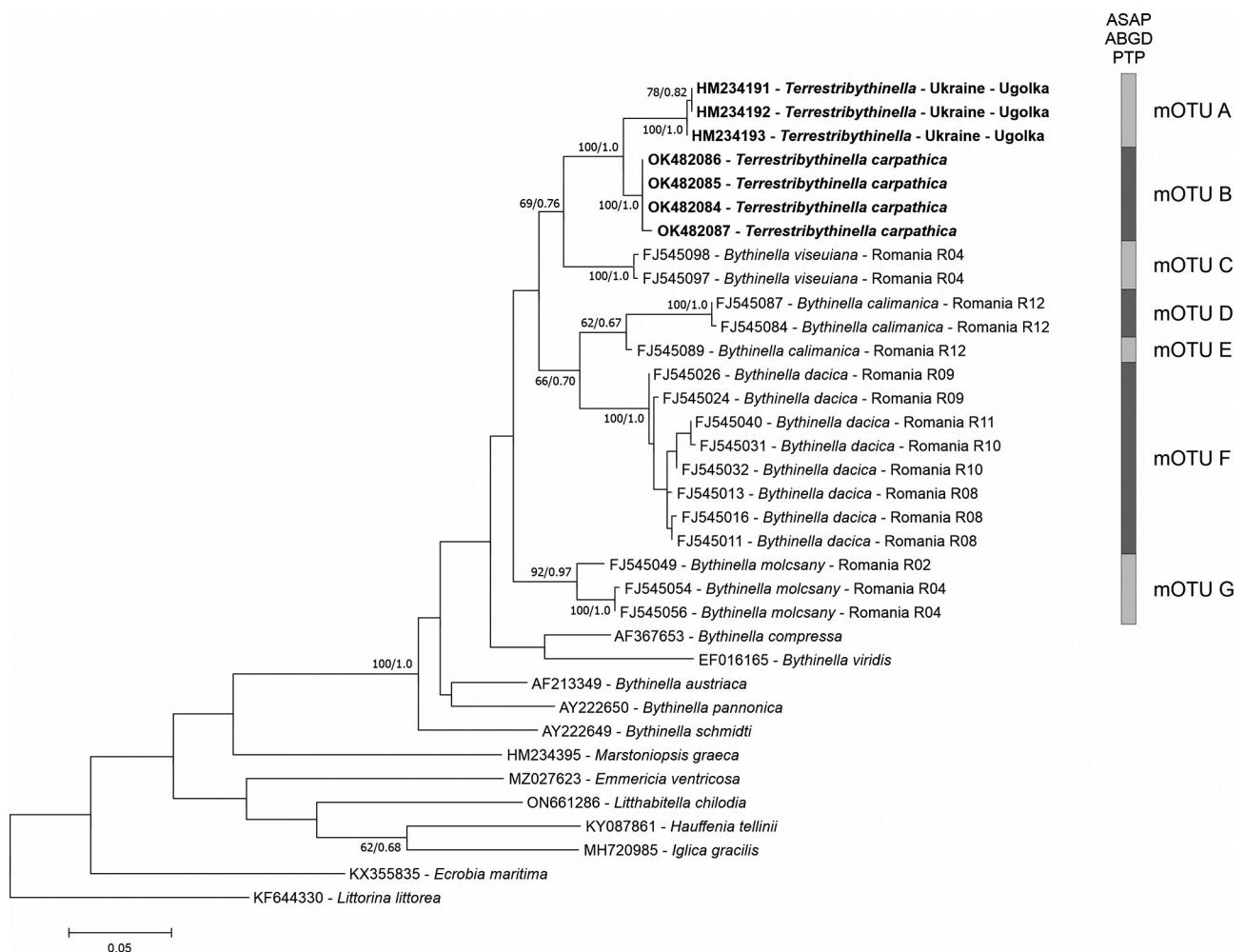


Fig. 1. Maximum likelihood tree based on the COI for *Terrestrabythinella* and related taxa, bootstrap supports (>60%) and Bayesian posterior probabilities are given. The HKY + I + G model were used for ML and BI analysis. Results of species delimitation are also shown. The GenBank numbers are given with the taxon, the sequences after WILKE et al. 2000, 2001, SZAROWSKA & WILKE 2004, BICHAIN et al. 2007a, FALNIOWSKI et al. 2009a, BENKE et al. 2011, ANISTRATENKO et al. 2021 (unpublished) and present paper

and close to the other *Bythinella* species (mOTUs D–G) from Romania, whose localities were not far from the Ukraine. Such close relationships rather definitely deny the possibility of such substantial anatomical differences as the ones listed above, described by SITNIKOVA et al. (1992). *Terrestrabythinella* clearly belongs to the genus *Bythinella*, and it cannot be accepted as a genus without creating a paraphyletic status of *Bythinella*. Thus, the only solution is to consider *Terrestrabythinella* a junior subjective synonym of *Bythinella* Moquin-Tandon, 1856. Phylogenetic analyses using all available *Bythinella* sequences also support this thesis (BENKE et al. 2011, JASZCZYNSKA in press).

Another genus reported in the Bythinellidae in WORMS (2024) is *Strandzhia* Georgiev et Glöer, 2013,

with its type species *S. bythinellopenia* Georgiev et Glöer, 2013. Its placement within the Bythinellidae remains enigmatic, since it was described as a member of the Hydrobiidae (GEORGIEV & GLÖER 2013). Our sequences of COI deposited in GenBank (PP752094-PP752096), as well as the one published by DELICADO et al. (2024: OP096318) undoubtedly classify *S. bythinellopenia* as belonging to the genus *Grossuana* Radoman, 1983, since in our tree *Strandzhia bythinellopenia* clusters within the mOTU A (Fig. 2), together with *Grossuana derventica* Georgiev et Glöer, 2013, *G. falniowskii* Georgiev, Glöer, Dedov et Irikov, 2015, and *G. thracica* Glöer et Georgiev 2009. The p-distance within this group does not exceed 0.001, and between this group (mOTU A) and other species of *Grossuana* (FALNIEWSKI et al. 2016)

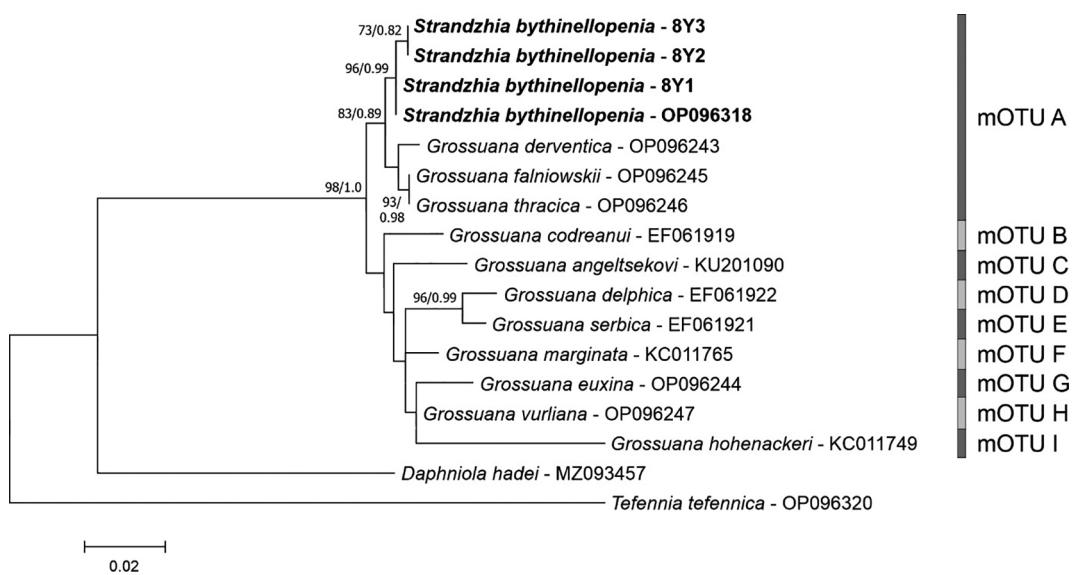


Fig. 2. Maximum likelihood tree based on the COI for *Strandzhia* and related taxa, bootstrap supports (>60%) and Bayesian posterior probabilities are given. The HKY+G model were used for ML and BI analysis. Results of species delimitation are also shown. The GenBank numbers are given with the taxon name, the sequences after SZAROWSKA et al. 2007, FALNIEWSKI et al. 2012, FALNIEWSKI et al. 2016, HOFMAN et al. 2021, DELICADO et al. 2024 and present paper

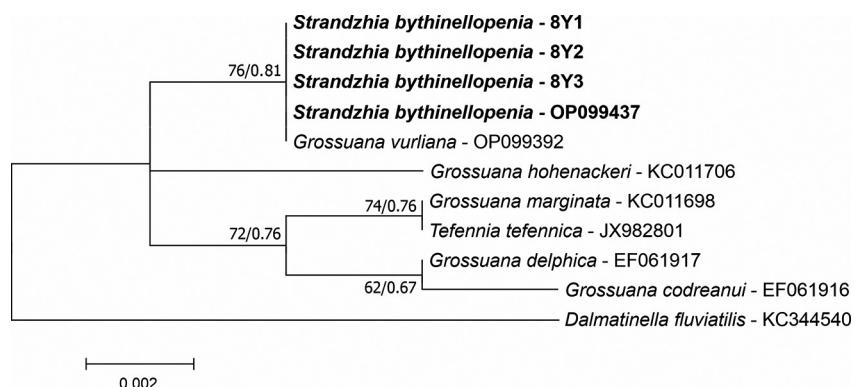


Fig. 3. Maximum likelihood tree based on the 18S for *Strandzhia* and related taxa, bootstrap supports (>60%) and Bayesian posterior probabilities are given. The HKY+G model were used for ML and BI analysis. The GenBank numbers are given with the taxon name, the sequences after SZAROWSKA et al. 2007, ÇAĞLAN et al. 2012, FALNIEWSKI et al. 2012, FALNIEWSKI & SZAROWSKA 2013, DELICADO et al. 2024 and present paper



varied from 0.013 to 0.064 (mOTUs B-I). The distance 0.001 is much below the threshold value generally accepted between congeneric species (e.g. PRIÉ & BICHAIN 2009, PRIÉ & CUCHERAT 2021). Also, 18S sequence (GenBank PP752097-PP752099) confirmed the assignment of *Strandzhia bythinellopenia* to the genus *Grossuana* (Fig. 3). Our dissection of paratypes of this species showed the simple penis typical of *Grossuana*. This, coupled with careful examination

of the photograph published by GEORGIEV & GLÖER (2013) clearly show that the region where the simple penis is bent typically for *Grossuana*, was erroneously interpreted as the base of bi-armed penis, and the base was misrecognised as the tip of the second arm. To conclude, *Strandzhia* Georgiev et Glöer, 2013 becomes a junior objective synonym of *Grossuana* Radoman, 1983.

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