New data on the valvatiform-shelled Hydrobiidae (Caenogastropoda, Truncatelloidea) from southern Greece

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Abstract
The minute valvatiform-shelled Hydrobiidae are less studied than other hydrobiid gastropods. In this paper, new data on these snails are presented, which have been collected at twelve springs in southern Greece: one in Boeotia, one on Evvoia Island, and ten on the Peloponnese Peninsula. Mitochondrial cytochrome oxidase subunit I (COI) and nuclear histone (H3) have been used to confirm the determinations and infer the relationships of the studied gastropods. They represent the genera Daphniola, Graecoarganiella and Isimerope. New localities, expanding the known geographic ranges, have been presented for Daphniola hadei and Daphniola louisi. A species of Daphniola found at two localities has been identified as a species new to science, and its description, including the shell, penis, and female reproductive organs is given. Possible relationships between Graecoarganiella and Isimerope are discussed; their representatives are possibly new species. At one locality a single specimen likely represents a new genus: it was found to be most closely related with Islamia, but genetically (p-distance) too distant to be congeneric with Islamia.

Keywords
COI, H3, molecular phylogeny, new species, Peloponnese, spring gastropods
**Introduction**

Minute freshwater gastropods with depressed trochiform (valvatiform) shells often were classified as belonging to the family Valvatidae Gray, 1840. The first genus described for such hydrobiid snails was *Horatia* Bourguignat, 1887 from Dalmatia (Schütt 1961; Radoman 1965; Szarowska and Falniowski 2014). Hydrobiidae in Greece are still poorly studied, and their microhabitats have drastically disappeared (Szarowska and Falniowski 2011a). The poor knowledge is perhaps of more concern for the valvatiform-shelled hydrobiids, since their low-spired tiny shells are easily overlooked or treated as juveniles. Some authors, for example Schütt (1980), have expanded the ranges of the Central European and North Balkan genera towards Greece, which was criticized by for example, Radoman (1983, 1985). In the present paper, we present the valvatiform-shelled gastropods collected at 12 localities in southern Greece.

**Material and methods**

The snails were collected by hand or with a sieve at twelve localities listed in Table 1 (Fig. 1) during two trips in 2009 and 2018. Samples were sieved through 500 μm sieve and fixed in 80% analytically pure ethanol, replaced twice, and sorted later. Next, the snails were put in fresh 80% analytically pure ethanol and kept at -20 °C in a refrigerator. The shells were photographed with a Canon EOS 50D digital camera, under a Nikon SMZ18 microscope with dark field. Dissections were performed under a Nikon SMZ18 microscope with dark field, equipped with Nikon DS-5 digital camera. Captured images were used to draw anatomical structures with a graphic tablet. Morphometric parameters of the shell were measured all by the same person using a Nikon DS-5 digital camera and ImageJ image analysis software (Rueden et al. 2017).

Snails for molecular analysis were fixed in 80% ethanol, changed twice, and later stored in 96% ethanol. DNA was extracted from whole specimens; tissues were hydrated in TE buffer (3 × 10 min); then total genomic DNA was extracted with the SHERLOCK extraction kit (A&A Biotechnology), and the final product was dissolved in 20 μl of tris-EDTA (TE) buffer. The extracted DNA was stored at -80 °C at the Department of Malacology, Institute of Zoology and Biomedical Research, Jagiellonian University in Kraków (Poland).

Mitochondrial cytochrome oxidase subunit I (COI), and nuclear histone 3 (H3) loci were sequenced. Details of PCR conditions, primers used and sequencing were given in Szarowska et al. (2016). Sequences were initially aligned in the MUSCLE (Edgar 2004) program in MEGA 7 (Kumar et al. 2016) and then checked in BIOEDIT 7.1.3.0 (Hall 1999). Uncorrected p-distances were calculated in MEGA 7. Estimation of the proportion of invariant sites and the saturation test (Xia 2000; Xia et al. 2003) were performed using DAMBE (Xia 2013). In the phylogenetic analysis, additional sequences from GenBank
Table 1. Geographic coordinates of identified sampling sites, by species. See also the map (Fig. 1). Extraction numbers (in bold) are also given, see also Figures 7, 8.

| Id | Site | Coordinates |
|----|------|-------------|
| 1  | W edge of Katarraktis, spring 564 m, Achaia, Peloponnese, Greece; **2A29, 2B24** | 38.1014, 21.8328 |
| 2  | Panagitsa, large spring 500 m, Arcadia, Peloponnese, Greece; **2A32, 2B26, 2B27** | 37.7725, 22.2219 |
| 3  | 3 km W of Megali Vrisi, Laconia, Peloponnese, Greece; **2A27** | 37.2267, 22.5222 |
| 4  | Spring beneath Katalfigio Parnonos, Laconia, Peloponnese, Greece; **2B19** | 37.2222, 22.6158 |
| 5  | Kastorio, spring, 3 3 km N of village at aquaducte, Laconia, Peloponnese, Greece; **2B20** | 37.1733, 22.2944 |
| 6  | Ag, Kiriaki spring N of Kato Kampia, Euboea, Greece; **2A33** | 38.5608, 23.8442 |
| 7  | Graecoarganiella parnassiana mouth of Erkinas Gorge, Kria 2, Boeotia, Livadia, Greece **2A28, 2B23** | 38.4319, 22.8750 |
| 8  | Isimerope semele Peloponnese, Achaia regional unit, Katarraktis center, spring and limestone cliffs, **2A30** | 38.0989, 21.8342 |
| 9  | Peloponnese, Achaia regional unit, Ag. Georgios (E of Tripotam), Vici spring, **2A31** | 37.8525, 21.9397 |
| 10 | Peloponnese, Achaia regional unit, Ag. Georgios (E of Tripotam), Anastasia spring, **2B21** | 37.8517, 21.9408 |
| 11 | Ladon spring E of Kerasia, Achaia regional unit, 474 m; 462 m alt., Greece, **2A22** | 37.8361, 22.1819 |
| cf. Islamia sp. 12 | Mili, spring below power station (on the Astros–Argos road), Argolis **2A34** | 37.5525, 22.7175 |

Figure 1. Localities of the sampling sites. For geographic coordinates see Table 1.
were used as references (Table 2). The data were analysed using approaches based on Bayesian Inference (BI) and Maximum Likelihood (ML). We applied the GTR model whose parameters were estimated by RAxML (Stamatakis 2014). In the BI analysis, the GTR + I + Γ model of nucleotide substitution was applied. The model was selected using MrModelTest 2.3 (Nylander 2004). The Bayesian analyses were run using MrBayes v. 3.2.3 (Ronquist et al. 2012) with defaults of most priors. Two simultaneous analyses were performed, each

### Table 2. Taxa used for phylogenetic analyses (COI and H3) with their GenBank (GB) accession numbers and references.

| Species                        | COI GB numbers | H3 GB numbers | References                                                                 |
|--------------------------------|----------------|---------------|-----------------------------------------------------------------------------|
| Agrafia wiktori Szarowska & Falniowski, 2011 | JF906762       | MG543158      | Szarowska and Falniowski 2011b; Grego et al. 2017                           |
| Alzoniella finalina Giusti & Bodon, 1984 | AF367650       | -             | Wilke et al. 2001                                                           |
| Anagastina zetwolvi (Radoman, 1973) | EF070616       | -             | Szarowska 2006                                                              |
| Belgrandiella kuesteri (Boeters, 1970) | MG551325       | MG551366      | Osiskowski et al. 2018                                                      |
| Dalmatinella fluviatilis Radoman, 1973 | KC344541       | -             | Falniowski and Szarowska 2013                                              |
| Daphniola dione Radea, Lampri, Bakolitsas & Parmakelis, 2021 | MW581160       | -             | Radea et al. 2021                                                           |
| Daphniola exigua (A. Schmidt, 1856) | EU047766, JF916470 | - | Falniowski et al. 2007; Falniowski and Szarowska 2011a                      |
| Daphniola gracilis Radoman, 1973 | EU047763       | -             | Falniowski et al. 2007                                                      |
| Daphniola haelei (Gittenberger, 1982) | JF916477, JF916479 | - | Falniowski and Szarowska 2011a                                              |
| Daphniola haelei (Gittenberger, 1982) | MZ093457, MZ093459, MZ265365– MZ265367 | present study | present study                                                              |
| Daphniola haelei (Gittenberger, 1982) | MZ093456, MZ265364 | - | Falniowski et al. 2007; Szarowska et al. 2014                               |
| Daphniola louisi Falniowski & Szarowska, 2000 | EU047769, KM887914, KM887915 | - | Falniowski et al. 2007; Szarowska et al. 2014                               |
| Daphniola louisi Falniowski & Szarowska, 2000 | MZ093456, MZ093460, MZ265368– MZ265372 | present study | present study                                                              |
| Daphniola magdalenar Falniowski, 2015 | KTT25578, KTT25580 | - | Falniowski and Sarbu 2015                                                   |
| Ecrobia maritima (Milaschewitsch, 1916) | KX355835, MG551322 | - | Osiskowski et al. 2016/Grego et al. 2017                                    |
| Fissuria bosi Boeters, 1981 | AF367654       | -             | Wilke et al. 2001                                                           |
| Graziana alpestris (Frauenfeld, 1863) | AF367641       | -             | Wilke et al. 2001                                                           |
| Graecoarganiella parnassiana Falniowski & Szarowska, 2011 | JN202349, JN202352 | - | Falniowski and Szarowska 2011b                                              |
| Graecoarganiella parnassiana Falniowski & Szarowska, 2011 | MZ093454, MZ093455, MZ265362– MZ265363 | present study | present study                                                              |
| Grossuana angelkowii Glöer & Georgiev, 2009 | KU201090       | -             | Falniowski et al. 2016                                                      |
| Grossuana bohnenckii (Küster, 1853) | KC011749       | -             | Falniowski et al. 2012                                                      |
| Hauffenia michleri (Kuščer, 1932) | KT236156, KY087878 | - | Falniowski and Szarowska 2015/Rysiewska et al. 2017                        |
| Isimerope semele Radea & Parmakelis, 2013 | KC841149       | -             | Radea et al. 2013                                                           |
| Isimerope semele Radea & Parmakelis, 2013 | MZ093450, MZ093453, MZ265355– MZ265361 | present study | present study                                                              |
| Isimerope sp. | JN202354       | -             | Falniowski and Szarowska 2011b                                              |
| Isimerope sp. | JN202354       | -             | Falniowski and Szarowska 2011b                                              |
| Islamia zermanica (Radoman, 1973) | KU662362, MG551320 | - | Beran et al. 2016; Grego et al. 2017                                       |
| Islamia sp. | MZ093465, MZ265373 | - | present study                                                              |
| Pontobelgrandiella sp. Radoman, 1978 | KG497024, MG551321 | - | Rysiewska et al. 2016/Grego et al. 2017                                   |
| Radomaniola curta (Küster, 1853) | KC011184       | -             | Falniowski et al. 2012                                                      |
| Sarajana apfelbecki (Brancsik, 1888) | MN031432, MN031438 | - | Hofman et al. 2019                                                          |
with 10,000,000 generations, with one cold chain and three heated chains, starting from random trees and sampling the trees every 1000 generations. The first 25% of the trees were discarded as burn-in. The analyses were summarised as a 50% majority-rule tree. Convergence was checked in Tracer v. 1.5 (Rambaut and Drummond 2009). FigTree v. 1.4.4 (Rambaut 2010) was used to visualise the trees. The ML analysis was conducted in RAxML v. 8.2.12 (Stamatakis 2014) using the RAxML-HPC v.8 on XSEDE (8.2.12) tool via the CIPRES Science Gateway (Miller et al. 2010). Bootstrap support was calculated with 1000 replicates and summarised on the best ML tree.

Abbreviations

GNHM  Goulandris Natural History Museum, Athens, Greece;
HNHM  Hungarian Natural History Museum, Budapest, Hungary;
JG    privet collection of Jozef Grego;
ZMUJ  Zoological Museum of the Jagiellonian University, Kraków, Poland;
ZPE   privet collection of Zoltán Péter Erőss.

Results and discussion

Systematics

Family Hydrobiidae Stimpson, 1865

Genus Daphniola Radoman, 1973

Notes. Radoman (1973) described this genus with its type species D. graeca Radoman, 1973, from the Daphne Spring in the valley of Tembe, North of Larissa. Schütt (1980) considered D. graeca a junior synonym of Valvata exigua Schmidt, 1856, described from “Greece”. Schütt (1980) designated a neotype from a group of small springs at Agia Paraskevi railway station, situated closely to the Daphne Spring, also in the valley of Tembe in Thessalia, but certainly not close to Thessaloniki as Kabat and Hershler (1993) state. Falniowski and Szarowska (2000) described Daphniola louisi from a small spring at the monastery at Kessariani, Athens, Attica. The description was not considered by Bodon et al. (2001), who followed either Schütt (1980) in synonymizing D. graeca with D. exigua, or Reischütz and Sattmann (1993) in including Valvata (Cincinna) hellenica Westerlund, 1898 in Daphniola exigua, thus rendering the genus Daphniola monotypic. Falniowski et al. (2007), applying soft-part morphology and anatomy as well as molecular markers, demonstrated the species distinctness of D. louisi, and identity of D. exigua with D. graeca. Gittenberger (1982) described Horatia hatei, a new species of Horatia he found 5 km SW of Yithion (Gythion), southern Peloponnese. Later, Falniowski and Szarowska (2011a) collected this gastropod close to the (probably) destroyed type locality, and both, morphology and molecular data confirmed its classification
within the genus *Daphniola*. Radea (2011) described *D. eptalophos* Radea, 2011 from the Parnassos Mountains. However, considering morphology, it is certainly not *Daphniola* especially in the intensively pigmented and massive penis. Its type locality is close (or rather the same) to the type locality of *Graecoarganiella parnassiana* Falniowski & Szarowska, 2011. Thus “*Daphniola etalophos*” is most probably a synonym of *Graecoarganiella parnassiana*, and clearly belongs to *Graecoarganiella* rather than *Daphniola*. Szarowska et al. (2014) found a few juvenile specimens (used for DNA sequencing) of *Daphniola* sp. on each of the two Aegean islands: Rhodos and Khios. Falniowski and Sarbu (2015) described *D. magdalenae* Falniowski, 2015 from the sulphide Melissotripa Cave in Thessalia. Finally, Radea et al. (2021) described *D. dione* Radea, Lampri, Bakolitsas & Parmakelis, 2021 from the Levkas Island (Ionian Sea), using morphology and molecular data. At two localities in North Peloponnese (1 and 2) we found another *Daphniola* taxon, whose morphology and COI sequences do not comply with any known *Daphniola* species.

*Daphniola longipenia* Grego & Falniowski, sp. nov.
http://zoobank.org/EE503BD0-65ED-4D53-AA37-ED75AB9448DA

**Types.** Ethanol-fixed specimens, Panagitsa, large spring, Arcadia, Peloponnese, Greece, 37°46′21″N, 22°13′19″E (Fig. 2), altitude 500 m, sieved from sand at the spring head; 26.09.2009; Z.P. Eröss, Z. Fehér, T. Fehér, J. Grego and A. Hunyadi coll., holotype: ZMUJ-M.2137; paratypes: ZMUJ-M.2138-ZMUJ-M.2139, two paratypes; HNHM-105279, 10 wet and 25 dry paratypes, GNHM 39591, 10 paratypes, ZPE 25 dry paratypes, JG F1198, 11 wet and 72 dry paratypes.

**GenBank numbers.** MZ093460–MZ093464; MZ265368–MZ265372

**Diagnosis.** Shell minute, valvatiform-trochiform, soft parts slightly pigmented, penis with extremely long and slender filament and small non-glandular outgrowth (lobe) on the left side. Readily distinguished from *D. exigua*, *D. louisi*, *D. magdalenae* and *D. dione* by the proportionally much lower spire of the shell, and the penis with a narrower base and a longer and thinner filament. Differentiated from the geographically (but not molecularly) most close *D. hadei* by the shell with usually lower spire, and the penis with smaller outgrowth and still longer and thinner filament.

**Description.** Shell (Fig. 3A–E) valvatiform-trochiform, up to 1.00 mm tall, having 3.5 whorls, spire height 10–12% height of shell. Apex flat. Teleoconch whorls moderately convex, evenly rounded, growing rapidly in diameter. Aperture slightly elliptical, parietal lip complete, umbilicus very broad, outer lip simple, orthocline. Teleoconch with delicate growth lines, periostracum pinkish or yellowish. Shell parameters for holotype and a series of paratypes are given in Table 3. Inner and outer sides of operculum smooth. Operculum pinkish. Animal brownish, with some spots of black pigment.

Female reproductive organs (Fig. 4) with a broad loop of the oviduct, a big bursa copulatrix with a long duct, and two moderately small receptacula seminis.
Figure 2. Type locality of *Daphniola longipenia* sp. nov. Panagitsa, Arcadia, Peloponnese A spring reservoir B spring head.

Figure 3. Shells of gastropods: *D. longipenia* A–D locality 2 (holotype, 2A32, 2B26, 2B27) E locality 1 (2A29); *Daphniola hadei* F locality 3 (2A27) G locality 4 (2B19) H locality 5 (2B20) I *D. louisi*, locality 6 (2A33) J–K *Graecoarganiella parnassiana*, locality 7 (2A28, 2B23) L–O *Isimerope* L locality 8 (2A30) M locality 9 (2A31) N locality 10 (2B21) O locality 11 (2A22) P cf. *Islamia* sp., locality 12 (2A34). Scale bar: 1 mm.
Table 3. Shell measurements of *Daphniola longipenia*; specimen symbols as in Figure 3; measured variables: see Figure 6.

|       | a     | b     | c     | d     | e     | a     |
|-------|-------|-------|-------|-------|-------|-------|
| A – holotype | 0.87  | 0.75  | 0.60  | 0.12  | 0.54  | 121   |
| B – 2A32  | 0.88  | 0.82  | 0.66  | 0.10  | 0.57  | 127   |
| C – 2B26  | 0.84  | 0.72  | 0.60  | 0.12  | 0.48  | 120   |
| D – 2B27  | 1.00  | 0.82  | 0.66  | 0.14  | 0.57  | 121   |
| E – 2A29  | 0.74  | 0.68  | 0.57  | 0.10  | 0.53  | 127   |
| M        | 0.866 | 0.758 | 0.618 | 0.116 | 0.538 | 123.200 |
| SD       | 0.093 | 0.062 | 0.040 | 0.017 | 0.037 | 3.493 |
| MIN      | 0.74  | 0.68  | 0.57  | 0.10  | 0.48  | 120   |
| MAX      | 1.00  | 0.82  | 0.66  | 0.14  | 0.57  | 127   |

Figure 4. Pallial and renal section of female reproductive organs of *Daphniola longipenia* [bc – bursa copulatrix, cbc – duct of bursa, ga – albuminoid gland, gn – nidamental gland, gp – gonoporus, ov – oviduct, ovl – loop of (renal) oviduct, rs – seminal receptacles (in black) rs\(_1\) and rs\(_2\) (as defined by Radoman 1973, 1983): rs\(_1\) – distal, rs\(_2\) – proximal]. Scale bar: 250 μm.

Penis (Fig. 5) extremely long and narrow, simple, with an almost vestigial outgrowth proximally on its left edge, and a prominent sharp-terminated filament, vas deferens not visible inside.

**Derivatio nominis.** The specific epithet *longipenia* refers to the extremely long filament of the penis.

**Distribution and habitat.** Apart from the type locality (our locality 2), this species was also found in the spring at the W edge of Katarraktis, Achaia, Peloponnese (our locality 1).

**Remarks.** Westerlund (1898) described *Valvata (Cincinna) hellenica* Westerlund, 1898, from “Vyteria in Arkadien”. Reischütz and Reischütz (2004) identified Westerlund’s “Vyteria” as Vitina, situated about 15 km from Panagitsa. They reported *Hauffenia hellenica* (Westerlund, 1898) also from Panagitsi spring. Our *D. longipenia* is most prob-
ably the gastropod reported by them. However, their identification of “Vyteria” as Vitina remains doubtful. The shell of the lectotype of *Valvata hellenica* presented by Reischütz and Sattmann (1993) looks different (Fig. 6) (enormously high and massive body whorl, another size and outline of the aperture). *Valvata hellenica* was reported several times from localities scattered throughout Greece, often in generic combination with *Hauffe-nia* or *Daphniola*. It can be assumed that these records report more than one species; or it was mentioned as a younger synonym under *Daphniola exigua* (e.g., Bodon et al. 2001). Summarising, the description of a new species is the most appropriate solution.

![Figure 5. Penis of *Daphniola longipenia*, bar equals 200 μm.](image)

![Figure 6. A shell of lectotype of *Valvata hellenica* presented by Reischütz and Sattmann (1993) B shell measurements: a – shell height, b – body whorl breadth, c – aperture height, d – spire height, e – aperture breadth, α – apex angle.](image)
In our trees (Figs 7, 8), *D. longipenia* is clearly distinct from all the other species of *Daphniola* and forms a distinct sister clade opposite to the remaining currently known *Daphniola* species. The high genetic distance (p-distance 0.106) can be found between *D. longipenia* and *D. hadei* (p-distance 0.106), whose localities are most close, and who share the most similar shell morphology, female reproductive organs and penis. In general, the genetic distances between *D. longipenia* and the other *Daphniola* varies from 0.097 (for *D. exigua*) to 0.141 (for *D. magdalena*) (Table 4).

*Daphniola hadei* (Gittenberger, 1982)

**GenBank numbers.** MZ093457–MZ093459; MZ265365–MZ265367

**Notes.** At the localities 3, 4 and 5 (Fig. 1, Table 1) gastropods were collected, whose shells (Fig. 3F–H), soft parts morphology and anatomy, clearly identified them as belonging to this species. Their molecular data (Figs 7, 8) were identical or nearly identical with the ones published in the GenBank. Their localities are situated somewhat north of the type locality.

*Daphniola louisi* Falniowski & Szarowska, 2000

**GenBank numbers.** MZ093456, MZ265364

**Notes.** The shell morphology (Fig. 3I), soft-part anatomy and molecular data (Figs 7, 8) of the gastropods collected at locality 6 (Fig. 1, Table 1), all showed that they belonged to this species. It has to be noted that the molecular variability in this species (p-distance 0.044) (Fig. 7) is much larger than in *D. hadei* (p-distance 0.013). The new record of *D. louisi*, located on Evvoia Island, considerably expands the range of the species, which so far was only known from Attica. The close phylogenetic relationship with the two juvenile *Daphniola* specimens collected on the Aegean islands, Rhodos and Khios, now combined with the present record from Evvoia, seems to confirm the ideas about the means of dispersal of *Daphniola* from continental Greece to the Aegean islands (Szarowska et al. 2014).

**Genus Graecoarganiella** Falniowski & Szarowska, 2011

**Graecoarganiella parnassiana** Falniowski & Szarowska, 2011

**GenBank numbers.** MZ093454–MZ093455; MZ265362–MZ265363

**Notes.** Falniowski and Szarowska (2011b) described a new, so far monotypic, genus of Hydrobiidae from Greece, Parnassus Mountains, S of Eptalofos, N of Kalania, found in a cistern and a small spring in a grassy pasture on a mountain pass. The type species, *G. parnassiana* Falniowski & Szarowska, 2011, is so far known only from the type locality. At the locality 7 (Fig. 1, Table 1), mouth of Erkinas Gorge, Kria 2,
Figure 7. Phylogenetic tree for COI showing relationships between the studied snails. Bootstrap supports (>60%) and Bayesian probabilities are given.
Table 4. P-distances for COI between main clades of the Daphniola.

|          | D. longipenia | D. hadei | D. dione | D. exiqua | D. magdalenae |
|----------|---------------|----------|----------|-----------|--------------|
| D. longipenia | 0.105       |          |          |           |              |
| D. hadei          | 0.088       | 0.080    |          |           |              |
| D. dione          | 0.097       | 0.092    | 0.087    |           |              |
| D. exiqua         | 0.141       | 0.153    | 0.133    | 0.154     |              |
| D. magdalenae     | 0.121       | 0.103    | 0.097    | 0.110     | 0.122        |

Boeotia, Livadia, we found gastropods, whose shells (Fig. 3J–K), and soft-part morphology indicated they belonged to Graecoarganiella, and were practically identical to G. parnassiana. Anatomy was not studied since the material was scarce and not fixed well enough. Our locality 7 is not far (about 35 km) from the type locality of G. parnassiana. The molecular data – partial sequences of COI – of our population showed
rather high distinctness (Fig. 7). However, as can be seen in the same phylogram, these differences (p-distance 0.038) are a little lower than the ones within *Daphniola louisi* (0.044). Thus, inclusion of our new population in *Graecoarganiella parnassiana* is seemingly justified.

**Genus *Isimerope* Radea & Parmakelis, 2013**

**Isimerope semele** Radea & Parmakelis, 2013

**GenBank numbers.** MZ093450–MZ093453; MZ265358–MZ265361

**Notes.** When describing *Graecoarganiella parnassiana* from the Parnassus Mts., Falniowski and Szarowska (2011b) reported three young hydrobiid specimens found at Mainalo Mountain, Peloponnesse, WSW of Piana, WNW of Tripolis, in a medium-sized spring and cistern. Their COI sequence was interpreted as indicating a distinct species congeneric with *Graecoarganiella parnassiana*. Later, Radea et al. (2013) found other species at Megali Vrisi, Pharmakas Mt., and described it as a representative of a new monotypic genus *Isimerope*, with *I. semele* as the type species. In our tree (Fig. 7) *Graecoarganiella* and *Isimerope* are quite distinct (p-distance 0.096), but form a well-supported clade (bootstrap value of 100%, Bayesian probability 1.0). The shells are very similar, and the same holds true for the radulae. The lack of a ctenidium, and egg capsules laid in the umbilicus of the shell, might be considered as unique shared character states. The penes and female reproductive organs of the compared taxa do not differ more than could be expected by different season of collection or fixation technique.

At the four localities: 8, 9, 10 and 11 (Fig. 1, Table 1) we collected gastropods, whose shells (Fig. 3L–O), soft parts morphology (not well-fixed material reduced the possible examination) and molecular data (Fig. 7) showed them as belonging to *Isimerope*. Again, as in the case of *Graecoarganiella*, our specimens of *Isimerope* may represent distinct species, but as in *Daphniola*, the molecular differences may be considered as within-species level variation (p-distance 0.035).

Our molecular data clearly show the close relationship of *Isimerope* and *Graecoarganiella*, contradicting their classification to different subfamilies (Belgrandiinae de Stefani, 1877 and Horatiinae D. W. Taylor, 1966, respectively), as stated in WORMS (WoRMS Editorial Board 2021). Both more anatomical and molecular data, as well as a broad-scale revision of the systematics of the Truncatelloidea proposed by Bouchet et al. (2017) are badly needed.

cf. *Islamia* sp.

**GenBank numbers.** MZ093465; MZ265373

**Notes.** At the locality 12, in Mili, Argolis, in a spring below the power station, a gastropod was found (Fig. 3P), whose molecularly inferred phylogenetic position (Fig. 8) remains enigmatic. Its sister taxon is *Islamia* Radoman, 1973. The clade’s bootstrap sup-
port for two concatenated loci is 100%, strongly suggesting that both mOTUs belong to the same taxon, but the genetic distance between them seems too high (p-distance 0.135). The p-distances in COI were 0.109 and 0.138 between this taxon and *Fissuria* Boeters, 1981 and *Alzoniella* Giusti & Bodon, 1984, respectively, although the shell morphology still suggests an affiliation with *Islamia*. Anyway, with only one more shell and lack of molecular data on the other Greek *Islamia* species, a justified taxonomic decision has to be postponed until more, and better, preserved specimens are available.

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