4-5-2012

A new species of springsnail (*Pyrgulopsis*) from the Owyhee River Basin, Nevada

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A NEW SPECIES OF SPRINGSNAIL (PYRGULOPSIS) FROM THE OWYHEE RIVER BASIN, NEVADA

Robert Hershler1,2 and Hsiu-Ping Liu3

ABSTRACT.—We describe a new springsnail species, Pyrgulopsis cybele, from the Owyhee River basin (northwestern Nevada) based on morphologic and molecular (mtCOI) evidence. Pyrgulopsis cybele differs from other members of its genus in its unique pattern of penial ornament, which consists of small glands on the distal edge of the penial lobe, base of the penial filament, and outer edge of the medial section of the penis. It is further differentiated from regional congeners by its thickened inner-shell lip and mtCOI sequences. A Bayesian analysis based on COI data placed P. cybele in a well-supported clade that contained congeners from Snake River, Great Basin, Colorado River, and California Pacific Coastal drainages; the sister taxon of this new species was not resolved. The COI divergence of P. cybele relative to its most genetically similar congener (P. glandulosa) suggests that it evolved subsequent to emplacement of the Miocene basalt that carpets the upper South Fork Owyhee basin. Pyrgulopsis cybele was collected from 2 closely proximal springs along the South Fork Owyhee River below the Nevada Pipeline Crossing. Although these populations are in a remote wilderness study area, they may be threatened by livestock grazing and human disturbance.

Pyrgulopsis is a western North American genus of hydrobiid gastropods that is distributed from the Snake–Columbia River basin and Missouri River headwaters to the lower Rio Grande basin (Hershler et al. 2008: fig. 1). The tiny, gill-breathing species in this genus typically live in springs and have very small geographic ranges (Hershler and Sada 2002). Pyrgulopsis contains 133 species (Hershler and Liu 2010, Hershler et al. 2010), more than half of which (78 of 133, 58%) have been described since 1995. The actual number of species in this genus is probably considerably larger than currently recognized, given that many putative novelties have not been formally treated taxonomically (e.g., Frest and Johannes 1995, Liu et al. 2003) and given that much of the West still has not been surveyed well for these tiny animals.

Four Pyrgulopsis species have been previously reported from the Owyhee River watershed, a remote and rugged subunit of the Snake River basin that occupies ca. 10,950 mi² (28,360 km²; Hardy et al. 2004) in Idaho, Nevada, and Oregon (Hershler 1998, Hershler and Liu 2009). Three of these species (Pyrgulopsis fresti, Pyrgulopsis intermedia, and Pyrgulopsis owyheensis) are distributed within a reach that extends from slightly above Three Forks to slightly above Owyhee Reservoir, Oregon; the fourth (Pyrgulopsis sadai) is found in the East Little Owyhee River drainage, Nevada (Fig. 1). In 2009 the first author discovered a population of Pyrgulopsis along the South Fork of the Owyhee River near the Oregon–Nevada border >100 km upflow from Three Forks (Fig. 1). A second, closely proximal population was discovered in
2010. Here we describe these populations as a new species of *Pyrgulopsis* based on morphological and molecular evidence. We also evaluate the phylogenetic relationships of this species and discuss its conservation status.

**METHODS**

Specimens were relaxed with menthol crystals and fixed in dilute formalin for anatomical study. Snails used for mtDNA sequencing were preserved in 90% ethanol in the field. Coordinates for the collecting localities were obtained in the field using a Garmin Oregon 450t GPS unit (UTM NAD 83). Types for the new species and other voucher material from this study were deposited in the Smithsonian Institution’s National Museum of Natural History (USNM) collection.

Five females and 5 males (all adults) from the type locality of the new species were dissected; large series of males from both localities were examined to assess variation in penial glandular ornament. Variation in the number of cusps on the radular teeth was assessed using the method of Hershler et al. (2007). Other methods of morphological study and descriptive terminology are those used in recent taxonomic
investigations of *Pyrgulopsis* (Hershler 1998, Hershler et al. 2003a). Basic statistics were compiled for the shell data using Systat for Windows 11.00.01 (SSI 2004).

Genomic DNA was extracted from entire snails using a CTAB protocol (Bucklin 1992). A 658 bp segment of cytochrome *c* oxidase subunit I (COI) corresponding to "Folmer's fragment" (Folmer et al. 1994) was amplified and sequenced with primers LCO1490 and HCOI2198 following protocols of Liu et al. (2003). Sequences were determined for both strands and then edited and aligned using Sequencherä version 4.8. We sequenced 3–6 specimens from the 2 samples of the new species described herein to assess variation. Sample information and GenBank accession numbers for the sequenced specimens utilized in this study are in Table 1. The new sequences reported herein were deposited in GenBank under accession numbers JN255350–JN255358 (Table 1).

Sequence divergences (uncorrected p distances) were calculated using MEGA5.05 (Tamura et al. 2011). Preliminary molecular phylogenetic analyses, which included most of the currently recognized species of *Pyrgulopsis*, consistently positioned the novelty described herein in a clade with 9 other congeners. The final analysis included all of these taxa and 2 congeners that belong to a different clade from the northern Lahontan basin; one of these congeners (*Pyrgulopsis gibba*) was used as the root.

**Phylogenetic relationships were inferred using Bayesian inference in MrBayes 3.12 (Ronquist and Huelsenbeck 2003). MrModeltest (Nylander 2004) selected the general time reversible model (GTR + G), which best fit the data under the Akaike information criterion. In the initial Bayesian analysis, the burn-in was set at 10% (10,000 generations) of the chain length (100,000 generations). Three runs were conducted in MrBayes using the General Time Reversible model (GTR + G) selected by MrModeltest and the default random tree option to determine when the log-likelihood sum reached a stable value (by plotting the log-likelihood scores of sample points against generation time). The ln likelihoods started around \(-7550\) and quickly converged upon a stable value of about \(-1700\) after 8000 generations. For the final run, Metropolis-coupled Markov chain Monte Carlo simulations were performed with 4 chains for 1,000,000 generations, and Markov chains were sampled at intervals of 10 generations to obtain 100,000 sample points. The sampled trees with branch lengths were used to generate a 50% majority-rule consensus tree with the first 5000 trees (equal to 50,000 generations) removed to ensure that the chain sampled a stationary portion.

**Systematic Description**

**Family Hydrobiidae Troschel, 1857**

**Subfamily Nymphophilinae Taylor, 1966**

**Genus Pyrgulopsis Call and Pilsbry, 1886**

**Type species:** *Pyrgula nevadensis* Stearns, 1883, by original designation.

**Diagnosis:** Liu and Hershler (2005:296).

**Pyrgulopsis cybele,** new species

**Types.—**Holotype, USNM 1148155, unnamed spring brook on east side of South Fork Owyhee River, Elko County, Nevada, N 4648124, E 524954, Zone 11, 1439 m elevation, coll. RH and Chris Hansen, 1 November 2010. Paratypes (from same lot), USNM 1157696 (ca. 400 specimens).

**ETYMOLOGY.—**This species is named after Cybele, the Roman goddess of nature.

**Referred material:** NEVADA. Elko County: USNM 1128557 (ca. 150 specimens), unnamed spring brook on east side of South Fork Owyhee River, ca. 100 m north of the above, N 4645315, E 525593, Zone 11, 1463 m elevation, coll. RH, 2 June 2009.

**Diagnosis.—**A small species of *Pyrgulopsis* having a trochoid to ovate-conic shell with highly convex whorls and a thickened inner apertural lip. Penis having a medium-sized lobe and medium-length filament; ornament consisting of penial gland, Dg1, and terminal gland.

**Description.—**Shell trochoid to ovate-conic (Fig. 2A–D); height 1.6–2.8 mm; whorls 3.25–4.00. Periostracum tan. Protoconch near planispiral, about 1.4 whors, diameter about 420 μm, initial portion weakly wrinkled (Fig. 2E, 2F). Teleoconch whors highly convex, shouldered, sutures impressed, last whorl sometimes slightly loosened behind aperture; smooth apart from collabral growth lines. Aperture large, ovate, angled above. Inner lip complete, usually slightly disjunct, rarely adnate, thickened internally, sometimes markedly so; columellar shelf usually absent, narrow to moderate width when present; outer lip usually thin, sometimes thickened adapically, prosocline. Umbilicus usually small, narrow, rarely absent. Shell measurements and
<table>
<thead>
<tr>
<th>Species</th>
<th>Specimen code</th>
<th>Locality</th>
<th>Accession number</th>
</tr>
</thead>
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<tr>
<td><em>cybele</em></td>
<td>P237A-F</td>
<td>Spring brook entering north side of South Fork Owyhee River, Elko Co., NV</td>
<td>JN255353–JN255358</td>
</tr>
<tr>
<td></td>
<td>P252A-C</td>
<td>Spring brook entering north side of South Fork Owyhee River, Elko Co., NV</td>
<td>JN255350–JN255352</td>
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<tr>
<td><em>gibba</em></td>
<td>P134B, D</td>
<td>Springs west of Fee Reservoir, Surprise Valley, Lassen Co., CA</td>
<td>AY197603(^a), AY246359(^b)</td>
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<tr>
<td><em>glandulosa</em></td>
<td></td>
<td>Nelson Place spring, Verde River basin, Yavapai Co., AZ</td>
<td>AY627959(^a)</td>
</tr>
<tr>
<td><em>imperialis</em></td>
<td>P140A, C</td>
<td>Spring, Thacker Pass, Kings River Valley, Humboldt Co., NV</td>
<td>AY379450(^c), AY246356(^b)</td>
</tr>
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<td><em>internedia</em></td>
<td>P1B, E</td>
<td>Crooked Creek, Highway 95 crossing, Owyhee River basin, Malheur Co., OR</td>
<td>AY379442(^b), AY246351(^b)</td>
</tr>
<tr>
<td></td>
<td>P2B, C</td>
<td>Crooked Creek at Crooked Creek State Wayside (19/v/2000), Owyhee River basin, Malheur Co., OR</td>
<td>AY246352(^b), AY246353(^b)</td>
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<td>P4B, C</td>
<td>Skylight Spring, Barren Valley, Malheur Co., OR</td>
<td>AY379444(^c), AY379445(^c)</td>
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<td>IP60A-D</td>
<td>Crooked Creek at Crooked Creek State Wayside (16/vii/2006), Owyhee River basin, Malheur Co., OR</td>
<td>FJ172460(^d), FJ172461(^d), FJ172462(^d), FJ172463(^d)</td>
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<td>IP67A, C</td>
<td>Spring tributary to Birch Creek, Owyhee River basin, Malheur Co., OR</td>
<td>FJ172468(^d), FJ172469(^d)</td>
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<td></td>
<td>P217A</td>
<td>Spring along Owyhee River, above Long Sweetwater Rapids, Owyhee River basin, Malheur Co., OR</td>
<td>FJ172488(^d)</td>
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<td>P222A, C</td>
<td>Spring on hillside, Jackson Hole, Owyhee River basin, Malheur Co., OR</td>
<td>FJ172496(^d), FJ172497(^d)</td>
</tr>
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<td>P223A, B</td>
<td>Mouth of Rinehart Creek, Owyhee River basin, Malheur Co., OR</td>
<td>FJ172498(^d), FJ172499(^d)</td>
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<td></td>
<td>P224C</td>
<td>Spring west of Two Mile Spring, Owyhee River basin, Malheur Co., OR</td>
<td>FJ172500(^d)</td>
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<td><em>militaris</em></td>
<td>P147A, C</td>
<td>Spring west of Soldier Meadow Ranch, Black Rock Desert, Humboldt Co., NV</td>
<td>AY197596(^a), AY246362(^b)</td>
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<tr>
<td><em>montezumensis</em></td>
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<td>Montezuma Well, Verde River basin, Yavapai Co., AZ</td>
<td>AY485552(^c)</td>
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<tr>
<td><em>morrisoni</em></td>
<td>Pnor5</td>
<td>Page Springs, Verde River basin, Yavapai Co., AZ</td>
<td>AY485551(^c)</td>
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<tr>
<td></td>
<td>P150A</td>
<td>Spring, Bubbling Pond Hatchery, Verde River basin, Yavapai Co., AZ</td>
<td>DQ364007(^f)</td>
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<td><em>robusta</em></td>
<td>D33C, D</td>
<td>XL Spring, Abert Lake basin, Lake Co., OR</td>
<td>AY326348(^b), AY246349(^b)</td>
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<td></td>
<td>P3B, D</td>
<td>Hughlet Spring, Harney basin, Harney Co., OR</td>
<td>AY379430(^c), AY379431(^c)</td>
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<td></td>
<td>P5B, C</td>
<td>South Fork Malheur Reservoir, Malheur Cave Road crossing, Malheur River basin, Harney Co., OR</td>
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<td>P179A, B</td>
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<td>AY379426(^c), AY379427(^c)</td>
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<td><em>simplex</em></td>
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<td>Spring near Strawberry, Verde River basin, Yavapai Co., AZ</td>
<td>AY627949(^g)</td>
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<td><em>sola</em></td>
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<td>Brown Springs, Verde River basin, Yavapai Co., AZ</td>
<td>AY627957(^g)</td>
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<td><em>stearnsiana</em></td>
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<td>Stream in Little Sycamore Canyon, southern California coastal drainage, Ventura Co., CA</td>
<td>QQ275094(^b)</td>
</tr>
</tbody>
</table>

\(^a\)Hershler et al. 2003b
\(^b\)Hershler and Liu 2004b
\(^c\)Hershler and Liu 2004a
\(^d\)Hershler and Liu 2009
\(^e\)Hurt 2004
\(^f\)Hershler and Liu 2008
\(^g\)Liu and Hershler 2005
\(^h\)Hershler and Liu 2010
Fig. 2. Scanning electron micrographs of shells and opercula of *Pyrgulopsis cybele*: A, holotype, USNM 1148155; B, shell, USNM 1157696; C and D, shells, USNM 1128557; E, shell apex, USNM 1157696; F close-up showing protoconch sculpture, USNM 1157696; G, operculum, outer side, USNM 1157696; H, operculum, inner side, USNM 1157696. Scale bars: A–D = 1.0 mm; E, F = 100 μm; G, H = 200 μm.
whorl counts for the holotype and paratypes are given in Table 2.

Operculum thin, amber, nuclear region and inner edge darker, multispiral with eccentric nucleus; last 0.5 whorl frilled on outer side (Fig. 2G); attachment scar border variably thickened (Fig. 2H). Radula (Fig. 3A) with about 58 well-formed rows of teeth. Central teeth about 20 μm wide, cutting edge highly concave (Fig. 3B); lateral cusps 5–7; central cusp rounded or weakly pointed, pointed, parallel-sided proximally; basal cusp 1, small; basal tongue rounded, about as long as lateral margins. Lateral tooth face rectangular, angled; central cusp pointed (Fig. 3C), sometimes bifurcate; lateral cusps 2–3 (inner), 3–5 (outer); outer wing medium width, variably flexed, about 200% length of cutting edge; basal tongue variably developed. Inner marginal teeth having 18–24 cusps (Fig. 3D). Outer marginal teeth having 20–31 cusps; inner edge with rectangular wing (Fig. 3E). Radular count data were from USNM 1157696.


Distribution and Habitat.—Pyrgulopsis cybele was collected from 2 small, closely proximal spring brooks (ca. 100 m apart) that discharge into a section of the South Fork Owyhee River known as the Devil’s Pinball (BLM 2011; Fig. 5). Neither of these springs is shown on the USGS 7.5-minute topographic map of this area (Rubber Hill, Nevada–Idaho Quadrangle, 1973). This area is ca. 5.25 km downflow from the Nevada Pipeline river crossing and ca. 50 km (airline) west-southwest of Owyhee, Nevada. Snails were moderately abundant on stones and cobble in the lower reaches of these spring brooks, which flow down steep talus slopes on the east side of the river. This species was not
found in several other springs that were surveyed during the first author’s kayak trip down the South Fork Owyhee River (from the Nevada Pipeline crossing to Three Forks) in June 2009.

Remarks.—This nymphophiline snail is assigned to *Pyrgulopsis* based on the superficial position of the bursa copulatrix and bursal duct on the albumen gland, presence of a single seminal receptacle, and diffuse pattern of mantle pigmentation (per Liu and Hershler 2005:296). *Pyrgulopsis cybele* is distinguished from all other congeners by its unique pattern of penial ornament consisting of small glandular units on the distal edge of lobe (terminal gland), base of the filament (penial gland), and outer edge of the medial section of the penis (Dg1; see Hershler and Sada 2002 for a discussion of penial variation in this genus). It is further distinguished from its Owyhee basin congeners by its thickened inner shell lip, and mtCOI sequences (see below). The 2 populations of *P. cybele* differed in shell size (Fig. 2A–2D), but were closely similar in all other morphologic details.

Fig. 3. Scanning electron micrographs of radula of *Pyrgulopsis cybele*, USNM 1157696: A, portion of radula ribbon; B, central teeth; C, lateral and inner marginal teeth; D, inner marginal tooth; E, outer marginal tooth. Scale bars = 10 μm.
Molecular Analysis

The Bayesian analysis (Fig. 6) delineated specimens of *P. cybele* as a well-supported (100% posterior probability) lineage nested within a (well-supported) clade composed of congeners from the Snake River, Great Basin, Colorado River, and California Pacific Coastal drainages. *Pyrgulopsis intermedia* and *P. robusta* formed a well-supported subgroup within this clade; the relationships of *P. cybele* and the other species were not well resolved. *Pyrgulopsis cybele* differed from other congeners included in this analysis by 3.0%–7.4% sequence divergence.

Five variable sites were detected among the 9 *Pyrgulopsis cybele* specimens that were sequenced, yielding 5 haplotypes (Table 3). These haplotypes differed from each other by 0.15%–0.46%. Two of the haplotypes (I, II) were shared by both populations of these species.
DISCUSSION

The morphologically distinctive novelty described herein does not appear to be closely related to *P. fresti*, which is endemic to springs along a lower reach of the Owyhee River (Hershler and Liu 2009), or to the 3 congener that are distributed in the Owyhee and other drainage basins. This finding is consistent with molecular evidence that regional *Pyrgulopsis* faunas typically are composites of phylogenetically diverse lineages (Liu and Hershler 2005). The springs inhabited by *P. cybele* discharge through Miocene basalt, which extensively carpets the South Fork Owyhee River watershed (Coats 1987, Foord et al. 1987: plate 1). The 3.0% ± 0.6% divergence of *P. cybele* relative to its most genetically similar congener (*P. glandulosa*) suggests that it evolved well subsequent to emplacement of this basalt, based on the COI molecular clock rate that has been established for *Pyrgulopsis* (1.62% per million years; Hershler and Liu 2008).

*Pyrgulopsis* species are highly vulnerable to extirpation owing to their typically small, fragile habitats and narrow geographic ranges (Melhop and Vaughn 1994, Melhop 1996). *Pyrgulopsis cybele* was collected in a large wilderness study area on public land (BLM 1991). The small streams inhabited by this snail appeared to be in good condition when visited by RH, although he did not visit their headspring sources. However, the local watershed has probably been impacted by the livestock grazing that has been occurring in this region since the late 1800s (this area is currently part of a large grazing allotment; NDA 2003). This reach of the South Fork Owyhee River watershed also may be impacted by human recreational activities (e.g., boating). The description of this locally endemic species contributes to a substantial body of evidence supporting recognition of the Owyhee Uplands as a distinct focal area of biodiversity (Vander Schaaf 1996).

ACKNOWLEDGMENTS

This project was supported (in part) by funds provided by the Center for Biological Diversity, Oregon Natural Desert Association (ONDA), and Western Watersheds Project. Brent Fenty (ONDA) was instrumental in helping to secure this funding. The Bureau of Land Management (Vale District) provided additional logistical support that was facilitated by Shaney Rockefeller. Chris Hansen, Dave Steele, and Dan
Thomas provided a large amount of assistance in the field. We thank 2 anonymous reviewers and WNAN editor Chris Walser for their constructive comments on a draft of this manuscript.

Table 3. Variation of COI (cytochrome c oxidase subunit I) within Pyrgulopsis cybele.

<table>
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<tr>
<th>Specimen</th>
<th>Haplotype</th>
<th>Base pair position</th>
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</thead>
<tbody>
<tr>
<td>P252A</td>
<td>I</td>
<td>G C C C C C</td>
</tr>
<tr>
<td>P252B</td>
<td>I</td>
<td>G C C C C C</td>
</tr>
<tr>
<td>P252C</td>
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<td>P237F</td>
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<td>G T C C C</td>
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</table>

Fig. 6. Bayesian tree based on the COI (cytochrome c oxidase subunit I) data set. Posterior probabilities for nodes are provided when the probabilities are ≥95%. Terminals are labeled as in Table 1.

LITERATURE CITED


FOLMER, O., M. BLACK, W. HOEH, R. LUTZ, AND R. VRIJENHOEK. 1994. DNA primers for amplification of


[Received 22 August 2011
Accepted 13 January 2012]