New Record of the Birdbeak Burrfish *Cyclichthys orbicularis* (Pisces: Diodontidae), in Korea

Won-Chan Lee¹, Jeong Bae Kim¹, Hyong Chul Kim¹, Seung Eun Bae², Jung-Hwa Ryu² and Jin-Koo Kim²*

¹National Fisheries Research and Development Institute, Busan 619-705, Korea  
²Department of Marine Biology, Pukyong National University, Busan 608-737, Korea  
³Ryujunghwa Marine Research Institute, Busan 614-811, Korea

**Abstract**

A total of six specimens (57.03-100.72 mm in standard length) belonging to the family Diodontidae were collected around Jeju Island, Korea between July and August 2013. On the Basis of morphological and molecular analyses, we identified the specimens as *Cyclichthys orbicularis* (Bloch, 1785), a new record to the Korean fish fauna. Morphologically, the species is characterized by nine caudal fin rays, immovable spines on the head and body except the caudal peduncle, and small black spots dorsally. Some morphometric characters changed disproportionately with growth, which was a new finding. Caudal peduncle length, body width and pre anus length to standard length tend to grow at a rate different from that of overall growth. When the 601 base pairs of mitochondrial DNA cytochrome c oxidase subunit I sequences of our specimens were compared with those of other diodontid fishes, our specimens clustered tightly around *C. orbicularis* (*d = 0.000-0.002*) in an NJ tree, with a high bootstrap value (99%). We herein describe the morphological and molecular traits of the specimens, and propose a new Korean name, “gud-ga-si-bok-sok” for the genus *Cyclichthys*, and “dung-geun-gud-ga-si-bok” for *C. orbicularis*.

**Key words:** *Cyclichthys orbicularis*, Diodontidae, Jeju Island, Korean fish fauna, New record

**Introduction**

The family Diodontidae of the order Tetraodontiformes comprises 19 species in 6 genera worldwide (Nelson, 2006), with 2 species in 2 genera in Korea (Kim et al., 2005) [*Diodon holocanthus* Linnaeus, 1958; *Chilomycterus affinis* (Linnaeus, 1758)]. Diodontid fishes are commonly known as porcupinefishes, burrfishes, and spiny puffers (Gallo et al., 2009). They are able to inflate the body, an important defense mechanism for such slow moving species (Cavaleri, 2000; Leis, 2006; Gallo et al., 2009). The body is covered with well-developed sharp spines, and species are identified according to whether the spines are movable or immovable (Leis, 1986; Aizawa, 2002; Leis, 2006). Some diodontid fishes are poisonous, having tetrodotoxin in their ovaries (females) and liver, and can also cause ciguatera (Allen and Erdmann, 2012; Bandyopadhyay, 2013). Most are generally scarce due to their solitary habits; they are found on soft bottoms, on rocky or coral reefs as well as coastal trawling grounds (Leis, 2001; Allen and Erdmann, 2012). Consequently, diodontid fishes have high bycatch rates by trawling and their recovery capacity is low (Stobutzki, 2001). Although Diodontidae fishes are not usually sold commercially for consumption, they are popular in the marine aquarium trade (Leis, 2001; Ellis, 2006). They are a conspicuous and readily captured species, and have been the focus of interest by naturalists for some time (Leis, 2006). However, in Korea, there have been few studies on diodontid fishes other than the brief description of the two species by
deposited in the ichthyology laboratory in Pukyong National University (PKU), Korea. Genomic DNA was extracted from muscle tissue using Chelex 100 resin (Bio-rad, USA), and then polymerase chain reaction (PCR) was performed using VF2 (5’-TCAACCAACCACAAAGACATTGGC-3’) and FishR1 (5’-TACAC TTCTGGGTGGCCAAAGAATCA-3’) primers that amplifies mtDNA COI (Ward et al., 2005; Ivanova et al., 2007). PCR was performed in a total volume of 30 μL which contained DNA template 1 μL, dNTP 2.4 μL, 10X buffer 3 μL, Taq polymerase 0.1 μL, reverse primer 1 μL, forward primer 1 μL, and distilled water, and conducted under the following conditions: initial denaturation was for 3 min at 94°C, followed by 35 cycles of 30 s at 94°C for denaturation, 30 s at 50°C for annealing, and 90 s at 72°C for extension, with a final extension at 72°C for 10 min. The PCR products were purified with Davinch™ PCR Purification Kit (Davinch-K, Seoul, Korea). The DNA was sequenced using an ABI 3730XL sequencer and an ABI PRISM® BigDye® Terminator v 3.0 Ready Reaction Cycle Sequencing Kit (Applied Biosystems, USA). The nucleotide sequence was deposited in the DDBJ/EMBL/GenBank databases (accession numbers: KJ626293‒KJ626296). The sequence was aligned with ClustalW (Thompson et al., 1994) in BioEdit version 7 (Hall, 1999). Sequences of four diodontid fishes (Cyclichthys orbicularis, C. spilostylus, Chilomycterus reticulatus, Diodon holocanthus), from the National Center for Biological Information database, were used for the comparison, and Takifugu niphobles was used as an outgroup. The genetic distances were calculated with the Kimura-2-parameter model (Kimura, 1980) in MEGA 5 (Tamura et al., 2011). An neighbor-joining

Material and Methods

A total of six specimens were collected around Jeju island, Korea between July and August 2013 (Fig. 1). Counts and measurements methods followed Nakabo (2002) with a Vernier caliper to the nearest 0.1 mm. Those specimens are

Table 1. Comparison of counts and measurements for Cyclichthys orbicularis and C. spilostylus

<table>
<thead>
<tr>
<th></th>
<th>Cyclichthys orbicularis</th>
<th>C. spilostylus</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>This study</td>
<td>Bloch (1785)</td>
</tr>
<tr>
<td>Number. of specimens</td>
<td>6</td>
<td>6</td>
</tr>
<tr>
<td>Standard length (mm)</td>
<td>57.03-100.72</td>
<td>100.4-144.4</td>
</tr>
<tr>
<td>Counts</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Dorsal fin rays</td>
<td>11-12</td>
<td>11</td>
</tr>
<tr>
<td>Pectoral fin rays</td>
<td>19-21</td>
<td>21</td>
</tr>
<tr>
<td>Anal fin rays</td>
<td>10-12</td>
<td>11</td>
</tr>
<tr>
<td>Caudal fin rays</td>
<td>9</td>
<td></td>
</tr>
<tr>
<td>Measurements</td>
<td></td>
<td></td>
</tr>
<tr>
<td>In % of standard length</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Head length</td>
<td>42.73-48.15 (44.89)</td>
<td></td>
</tr>
<tr>
<td>Caudal peduncle length</td>
<td>8.64-12.69 (10.74)</td>
<td>-</td>
</tr>
<tr>
<td>Caudal peduncle depth</td>
<td>5.90-6.96 (6.48)</td>
<td>-</td>
</tr>
<tr>
<td>Eye diameter</td>
<td>9.57-11.83 (10.49)</td>
<td>-</td>
</tr>
<tr>
<td>Head width</td>
<td>35.65-40.74 (38.56)</td>
<td>-</td>
</tr>
<tr>
<td>Body width</td>
<td>53.54-72.49 (63.06)</td>
<td>-</td>
</tr>
<tr>
<td>Interorbital width</td>
<td>26.44-28.70 (27.27)</td>
<td>-</td>
</tr>
<tr>
<td>Nostril to mouth</td>
<td>9.22-12.55 (10.92)</td>
<td>-</td>
</tr>
<tr>
<td>Height of gill opening</td>
<td>9.82-12.13 (11.02)</td>
<td>-</td>
</tr>
<tr>
<td>Mouth width</td>
<td>13.86-16.00 (15.09)</td>
<td>-</td>
</tr>
<tr>
<td>Predorsal length</td>
<td>80.44-90.76 (86.06)</td>
<td>-</td>
</tr>
<tr>
<td>Preanal length</td>
<td>84.15-93.86 (89.30)</td>
<td>-</td>
</tr>
</tbody>
</table>

Fig. 1. Sampling area of Cyclichthys orbicularis in the present study.
Description

Spines of body and head with 3 bases, except in *C. spilostylus* which has some spines on top of head with 4 bases; no spines on caudal peduncle; 9 caudal fin rays; no spots on fins. The most similar genus, *Chilomycterus*, is distinguishable from *Cyclichthys* in having a small spine on dorsal side of caudal peduncle, 10 caudal fin rays, and spots on fins (Leis, 1986).

*Cyclichthys orbicularis* (Bloch, 1785) (Fig. 2)
(New Korean species name: Dung-geun-gud-ga-si-bok)

*Diodon orbicularis* Bloch 1785: 73 (Type locality: Jamaica); Allen & Swainston, 1988: 156, Australia; Matsuura & Sakai, 1993: 372, Japan; Mishra & Krishnam, 2003: 51, India.

*Chilomycterus paracomaculatus* von Bonde, 1923: 38 (Type locality: Union of South Africa).


Distribution

Found in the tropic waters of the Indo-West Pacific (Leis, 2001) and Southeast Atlantic (Leis, 1986) Oceans, near reef areas, at depths of 9-170 m (Sommer et al., 1996).

MiDNA COI sequences analysis

Comparison of the 601 bp of mtDNA COI sequences of four specimens with those of other diodontid fishes revealed that the sequences of the specimens aligned best with that of *C. orbicularis* ($d = 0.000-0.002$), but clearly differed from those of *C. spilostylus* ($d = 0.066-0.068$), *Chilomycterus reticulatus* ($d = 0.105-0.108$), and *Diodon holocanthus* ($d = 0.105-0.107$). In an NJ tree, the specimens clustered with *C. orbicularis* (GU804950), with a high bootstrap value of 99% (Fig. 3).
Remarks

Our specimens collected around Jeju Island are placed in the genus *Cyclichthys* because they possess immovable spines with 3-rooted bases, 9 caudal fin rays, no spots on the fins, and no spines on the caudal peduncle (Leis, 1986, 2001). The specimens also matched the original description of *C. orbicularis* in most meristic counts, the presence of black spots in clusters dorsally and dorsolaterally, and three spines on top of eyes (Bloch, 1785; Leis, 2006; Allen and Erdmann, 2012). Molecular analysis supported the identification as *C. orbicularis* ($d = 0.000-0.002$).

*Cyclichthys orbicularis* differs from *Chilomycterus reticulatus* in the number of caudal fin rays (9 in *Cyclichthys orbicularis* vs. 10 in *Chilomycterus reticulatus*), the spine on dorsal surface of caudal peduncle (absent vs. present), and black spots on all fins (absent vs. present) (Leis, 1986; Aizawa, 2002; Kim et al., 2005). Although, according to Matsuura et al. (1993), several measurements of *Cyclichthys orbicularis* overlap with those of the morphologically similar *C. spilostylus*, the former is well distinguished from the latter by the number of subdermal roots (all spines with 3 roots in *C. orbicularis* vs. some spines with 4 roots in *C. spilostylus*), and clusters of dark spots on ventral side of body (absent vs. present) (Leis, 2001; Aizawa, 2002). Most measurements of the specimens in this study overlapped those of *C. orbicularis* in Japan, but the specimens showed significant differences in body width, caudal peduncle length, and preanus length (Table 1). Generally, body shape in fishes is associated with a variety of environmental factors, type of food or feeding mode, and sexual differences as well as growth period (Marcil et al., 2006; Frederich et al., 2008; Fruciano et al., 2011; Song et al., 2013). But, when standard length was regressed against the various measurements, caudal peduncle length showed positive correlation (Fig. 4a), whereas body width and preanus length were negatively correlated with standard length (Fig. 4b, 4c). Therefore, our results suggested that body shape changes with growth, not due to regional variation. We pro-

**Fig. 3.** Neighbor-joining (NJ) tree for cytochrome oxidase subunit I (COI) gene sequences of 4 diodontid fishes. The NJ tree was constructed under the K2P model using *Takifugu niphobles* as the outgroup. Superscripts indicate the registration number of voucher specimens.

**Fig. 4.** Comparison of relationships between standard length (SL) and proportional measurements of caudal peduncle length (A), body width (B), and preanus length (C) for *Cyclichthys orbicularis* from Korea (rectangle) and from Japan (diamond).
pose the new Korean name “gud-ga-si-bok-sok” for the genus *Cyclichthys*, and “dung-geun-gud-ga-si-bok” for *C. orbicularis* based on the species’ rounded shape.

**Acknowledgement**

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**References**


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