

Barbus ida, a new barbel species from the Southern Marmara Sea basin (Teleostei: Cyprinidae)

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Barbus ida, a new species, is described from the streams Gönen and Biga, in western Anatolia. It is distinguished from other *Barbus* species in the adjacent basins by a less thickened last unbranched dorsal-fin ray, serrated along the proximal half of its posterior margin; larger irregular black blotches on the back and the flanks, and small black spots on the head, extending downwards to the cheeks; 56–61 total lateral line scales, 11–13 scale rows between dorsal-fin origin and lateral line and 7–8 scale rows between anal-fin origin and lateral line. *Barbus ida* also differs from its most closely related species, *B. niluferensis*, by 8 nucleotide substitution sites in the mtDNA cytochrome oxidase I barcode region.

Introduction

A total of 11 species of the cyprinid genus *Barbus* have been reported from Turkish inland waters (Güçlü et al., 2020). Eight of which have been reported from Western Anatolia and the Thrace region, including: *B. anatolicus* Turan, Kaya, Geiger & Freyhof 2018 from Yeşilırmak and Kızılırmak rivers, *B. cyclolepis* Heckel, 1837 from Ergene River and İstranca stream, *B. escherichii* Steindachner, 1897 from Sakarya River, *B. niluferensis* Turan, Kottelat & Ekmekci, 2009 from Susurluk River, *B. oligolepis* Battalgil, 1941 from the streams and rivers in Biga Peninsula, *B. xanthos* Güçlü, Kalaycı, Küçük & Turan, 2020 and *B. pergamonicus* Kara-

man, 1971 from the streams and rivers in the Aegean Sea basin, and *B. tauricus* Kessler, 1877 from small streams in southeastern and western Black Sea (Kottelat & Freyhof, 2007; Geiger et al., 2014; Turan et al., 2018; Levin et al., 2019; Güçlü et al., 2020).

Geiger et al. (2014) recorded *Barbus* (as *Barbus* sp.) from Gönen Stream. Later on, we examined additional specimens of *Barbus* sp. from both Gönen and Biga streams and compared these with the morphologically similar (and putatively genetically similar) species, *B. niluferensis*, and also species in adjacent basin. Based on morphological and genetic characters, we consider the specimens of

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