

9th Conference of the Scandinavian - Baltic Society for Parasitology

PARASITES IN A CHANGING WORLD

ABSTRACT BOOK



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Capturing boundaries in a group of *Parvatrema* (Digenea:Gymnophallidae) cryptic species

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Within the digenean genus Parvatrema (Gymnophallidae) there is a group of species that share a unique feature: parthenogenetic metacercariae in the molluscan second intermediate host (2IH). We found that such species are common at the sea shores in the northern Holartic, and they use several gastropod species as 2IH. Species of Parvatrema differ in their interactions with 2IH: from commensalism in the extrapallial cavity to parasitism in the hepatopancreas and the gonad. Distinguishing between the metacercariae of different species is complicated because morphologically they are almost identical. rDNAbased genetic analysis also did not provide understanding of relationships between the Parvatrema isolates from different 2IH. We sequenced and annotated the mitochondrial genome of Parvatrema sp. (Cercaria quadriramis) from Littorina saxatilis, and used it for primer design. We then sequenced the cox1 gene fragment for all available Parvatrema isolates (23). The resulting alignment was analyzed with ABGD (https://bioinfo.mnhn.fr/abi/public/abgd/), and distribution of pairwise distances indicated the barcode gap presence and delimitation of five species: Cercaria quadriramis, Parvatrema homoeotecnum forma A, Parvatrema margaritense, Cercaria falsicingula, Parvatrema sp. 1. Phylogenetic analysis showed strict geographic attribution of these species either to the North Atlantic (NA), or to the North Pacific (NP). Presumably, the expansion of *Parvatrema* spp. proceeded from NP to NA, with two independent transfer events. Morphological similarity of the studied *Parvatrema* spp. suggests that they can be considered cryptic species that diverged as they discovered new 2IH species and new geographic region (NA). The research was funded by the RSF grant #18-14-00170.