

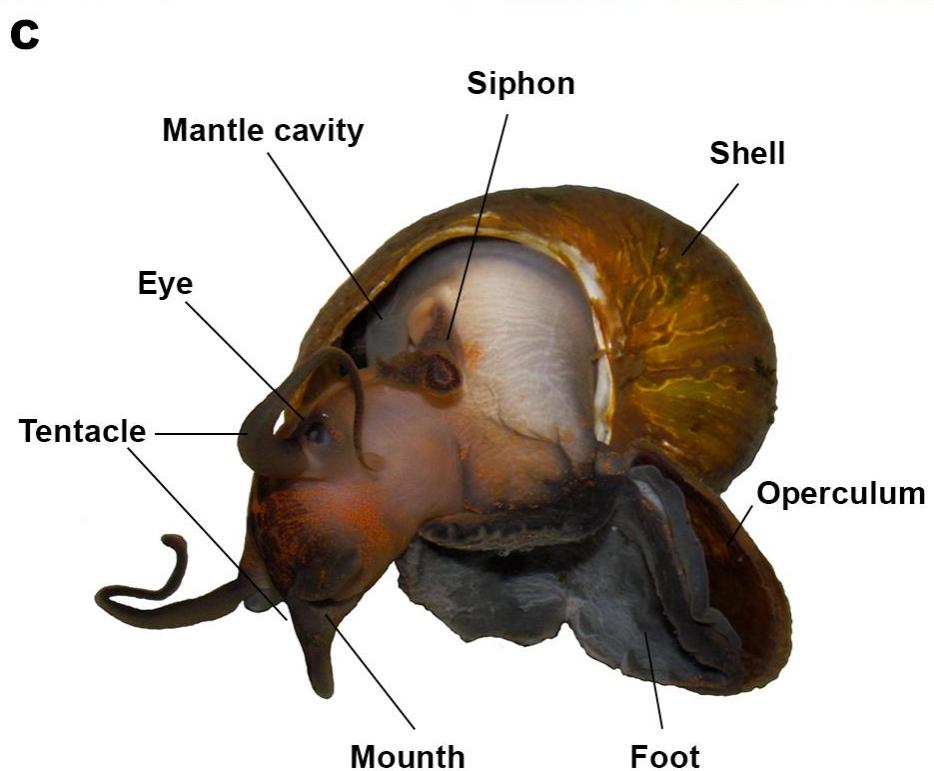
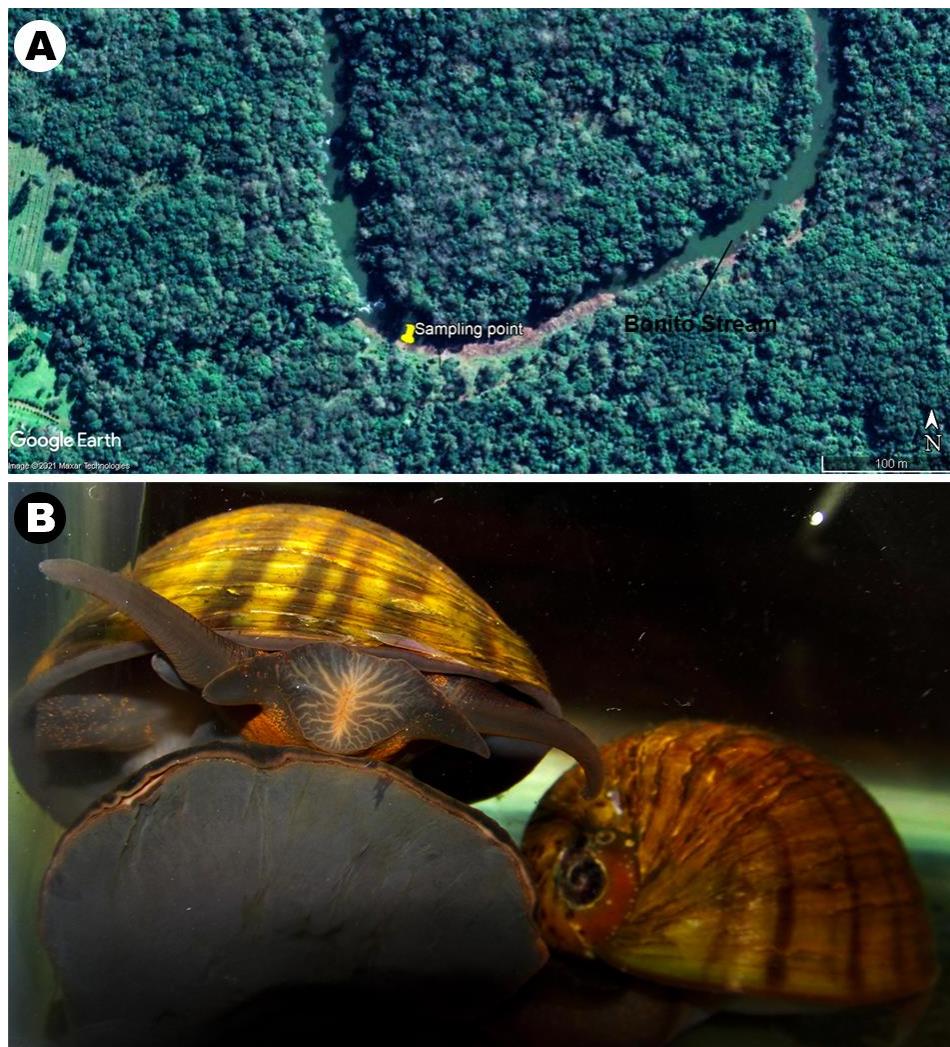
First morphological and molecular identification of the cercaria of *Stomylotrema vicarium*  
from the endemic apple snail *Pomacea americanista*

Supplementary Material

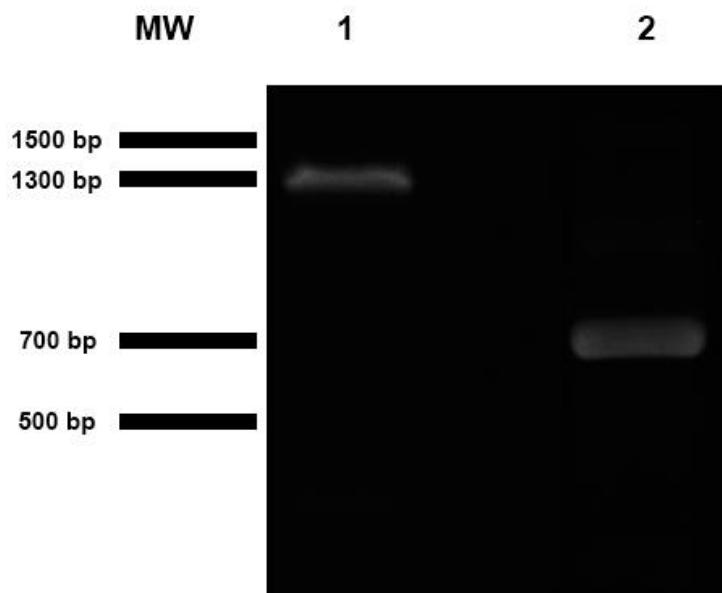
Federico A. Dellagnola<sup>1,2,3</sup>, Alejandra D. Campoy-Díaz<sup>1,2</sup>, and Israel A. Vega<sup>1,2,3,\*</sup>

<sup>1</sup>IHEM, Universidad Nacional de Cuyo, CONICET; <sup>2</sup>Universidad Nacional de Cuyo, Facultad de Ciencias Médicas, Instituto de Fisiología; <sup>3</sup>Universidad Nacional de Cuyo and Facultad de Ciencias Exactas y Naturales, Departamento de Biología

**\*Author for correspondence:** Israel A. Vega, [ivega@mendoza-conicet.gob.ar](mailto:ivega@mendoza-conicet.gob.ar)



**Supplementary Figure 1. The host *Pomacea americanista*.** **A.** Satellite view (Google Earth Inc.) from the specimen's sample sites (Bonito Stream, Misiones rainforest, Argentina). **B.** Alive specimens of *Pomacea americanista* in the aquarium. **C.** External morphology of a live specimen.



**Supplementary Figure 2.** Representative agarose gel (1.2 %) containing the PCR amplification products for the 28S rDNA (lane 1) and ITS1 (lane 2). Total DNA from intramolluscan stages isolated from the digestive gland of *P. americanista* was used as template. Lane 1: amplicon of~1300 bp that includes a whole region of the gene that encodes for the 28S rRNA. Lane 2: amplicon of~700 bp that includes the complete non-coding region of the Internal Transcribed Spacer (ITS1) placed on between the 18S rDNA and 5.8S rDNA genes.

**Supplementary Figure 3.** Trematode sequences isolated from the apple snail *P. americanista*

**A. rRNA 28S sequence**

>MW480895 *Stomylotrema vicarium* isolate DG\_Pam01 large subunit ribosomal RNA gene, partial sequence.  
TGAACAGGGACAAGCCCAGCACCAGAACCTGGCATGGCAATGGGTTTCAGGTGATCCGGTA  
GTCACTGCTCCACCCCTAAGTCATTAAATGAGTACGGTATTATGGACATGGCCGAAGAGGGTGAAAGGCCGTGG  
GGTGGAGTGCAGGTAGGCCAGTGAAGTGCCTGGAAGGACCTTGGAGTCGGTTGTTGAAATGCAGCCAAAGTGG  
GTGGTAAACTCCATCCAAGGCTAAATACAAGCACGAGTCCGATAGCGAACAGTACCGTGAGGGAAAGTTGAAAA  
GTACTTTGAAGAGAGAGTAAACAGTGCCTGAAACCGGTAGAGGTAAGCAGGTGGAGTTGAACGTCAAGCTCTGG  
GAATTCAACTGGTGGAGATTGGTTGAGCTGGTTGAATCGGTTGGCCTGAAAGTCTGTAGCAGCAGGCTCTGT  
TTTGACAGAGGTGCGCGATACACTTGTCAAGTGTGCGCTTCGGGTGCTCCGGACCGGCTGCCAGTGCAC  
TTCTCAGAGTGGTCACCACCGACCGGACTGCTGTGGCTGTTGCGGTTAACCGGCTTGCATTGTCCTGTGGCTC  
TGCTTGCCTGGGATGGCAGGTAGCCGTTGGCTTGTAGCTGCTACAGGCTTCGGTCTCCGGGTGTAATCAG  
CTGACCGCATCAGTCTGTGCAAGCGTCTGGAGACGGCGCTCGTGCCTGCGTGCCTGCGCTGGCGTGC  
TTCGGGTTGGTCACTATGTTGCCTGTTGGCAGGGCTGGTGATGGCCGGCTCGTCCGGTCCGGTGGCGTGC  
GTGGCGCGAAAGCAGGGGCCATAGTCTGTGGTAGTGGTAGACTATCCACCTGACCCGTCTGAAACACGGACC  
AAGGAGAGTAACATGTGCGCAAGTCATTGGCACTACGAAACCTATAGGCAAAGTGAAGGTCTGGCTTGT  
CCAGACTGAGGTGGATCCTGCTGTTCTCATGCGTGGTACCGCCAAGCTCGAGCGGAGGGCGCAGCACGGCCC  
GTCGGATGACAATGTCCAACGGCAATAACCAAGTCGGGCGGAGCATGAGCGTACATGTTGAGACCCGAAAGATG  
GTGAACATATGTTGCGCAGGTTGAAGCCAGAGGAAACTCTGGTGGAGGACCGAAGCGGTTGACGTGCAAATCGA  
TCGCCAGACGTGAGTATAGGGCGAAAGACTAATCGAAC

**B rRNA 18S-ITS 1-rRNA 5.8S sequence**

>MW481318 *Stomylotrema vicarium* isolate DG\_Pam01 small subunit ribosomal RNA gene, partial sequence; internal transcribed spacer 1, complete sequence; and 5.8S ribosomal RNA gene, partial sequence.

TGGTAAGTGCAGTCATAAGCTTGCCTGATTACGTCCTGCCCTTGTACACAYCGCCCGTCTACTACCGATTG  
AATGGTTAGCAAGGTCTCGGATTGGCGTATTGTATTGCTTCGGCAGCTGACCCGGTGTGAGAAGCCGACGAA  
TCTTGATATTGGAGGAAGTGGAAAGTCGTAACCACATTCCGTAGGTGAAACCTGCGGAAGAACGATTACCGTATT  
CCCCCTCCGAAAGCTGTACCTGGCTTAATGGGCCTACGTACAGCTTCTACTCTATTCCGGTCTTACCCCTCT  
AAAGCTATCGGGGTGCTTGTACCTTGTGGTAGTCAGTCCACTCTGGAAAGTGAAGGTTGTGCTGACCAAATCAGTG  
CTAGGCTTAATGAATGGTGTCTCAGCTACGGCTAGCCCCTAGCTTCCCTGTTGTAAAACCGTTAAAG  
TGGTACATCTGTGCCATTGCTCAACATGCACCCGGTGTGAACGGACTGCATGTGCAGTCGCTGGCGGTGCCTTA  
TCCCGGGCTGGACTGAGAAACCATTGGGTGTTCAAGGAAACTGAATGCTCAGTGTAGTACAACACTGATCGGTGGAT  
CACTCGGCTCGTGTGCGATGAAGAGCGCAGC