

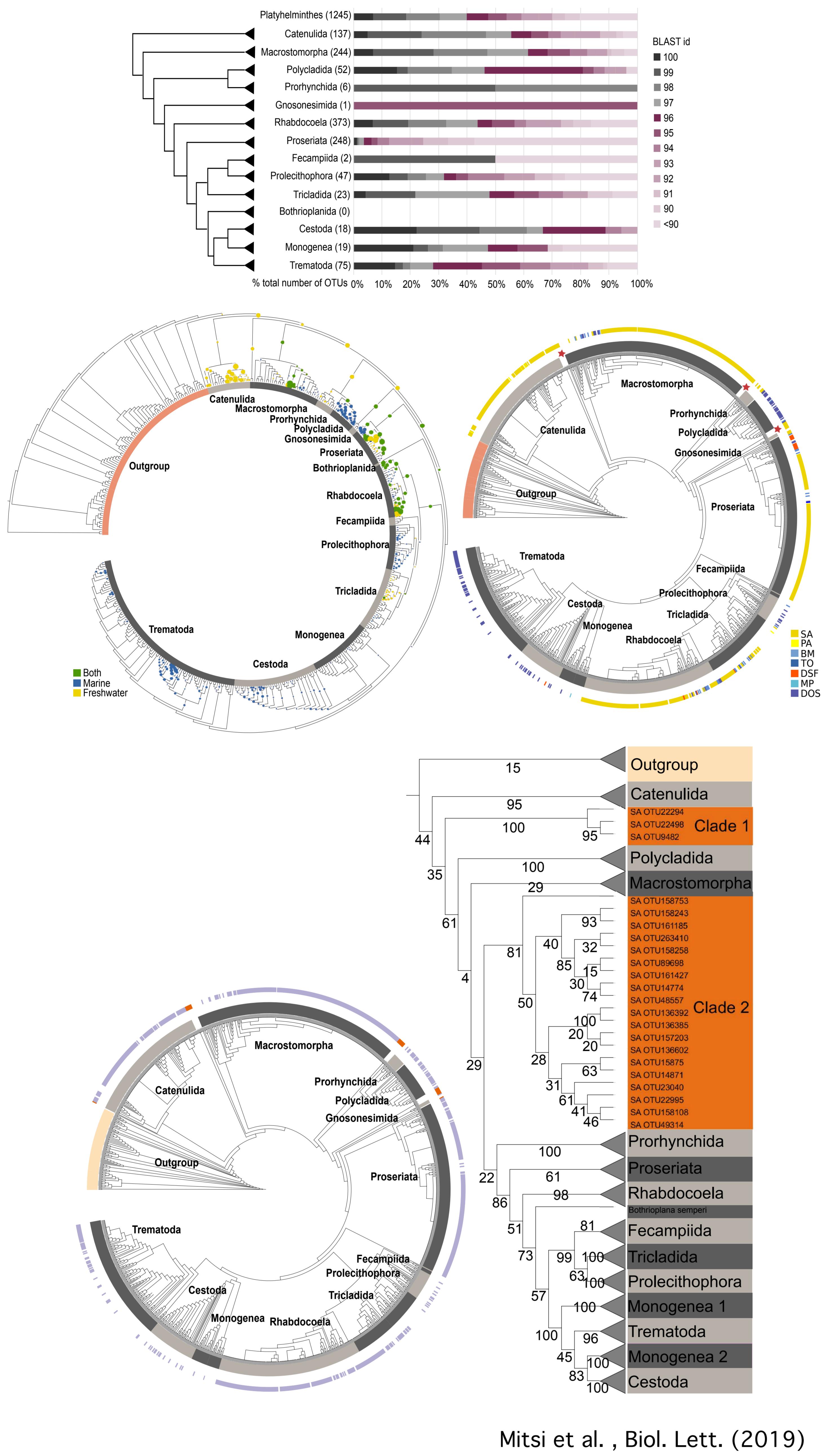


Eukaryotic diversity through the lens of metabarcoding and metagenomics

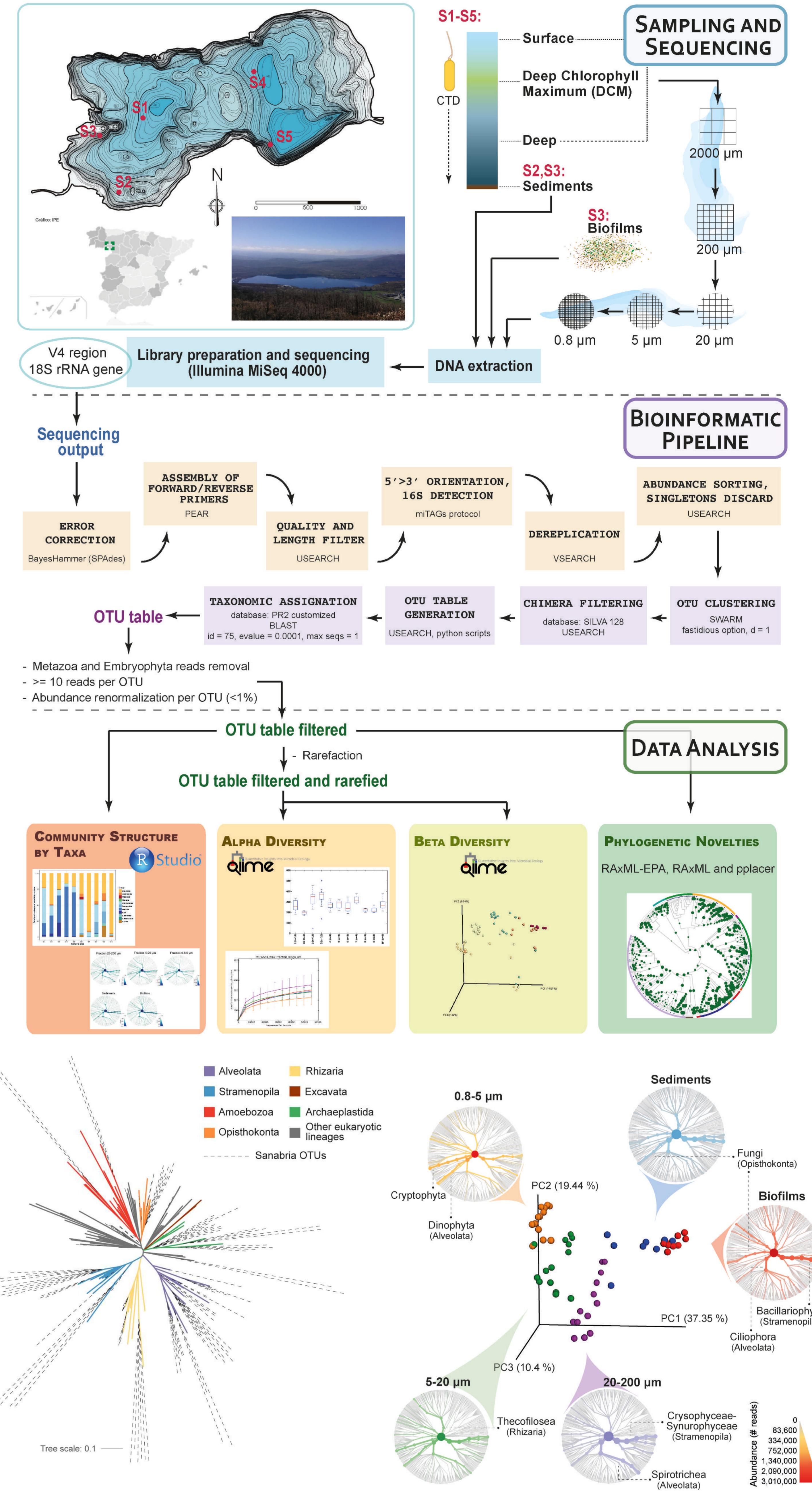
Konstantina Mitsi - Prof. Iñaki Ruiz-Trillo



P1. A global metabarcoding analysis expands molecular diversity of Platyhelminthes and reveals novel early-branching clades



P2. Assessing eukaryotic diversity in a glacial lake under eutrophication : the case study of Lake Sanabria



P3. The genome of *Txikispora* sp., the first filasterean parasite

