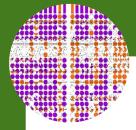
MOLECULAR BIOLOGY OF PROTISTS



Explore the evolution of eukaryotic genomes. We are studying the gene repertoire evolution in Euglenozoa, a group of unicellular eukaryotes that includes human pathogens of the genera *Trypanosoma* and *Leishmania*, as well as their well-known free-living relatives such as *Euglena gracilis* and diplonemids.

Our research focuses on investigating the genetic changes that underlie lifestyle switches and the genes that define the ecological success of these organisms. We also study the evolution of non-coding DNA. For that we employ cutting-edge methods of bioinformatics and a combination of omics data, including genomic, transcriptomic, proteomic, and metabolomic data.



Join our research team!

I am an experienced bioinformatician (more than 10 years of experience) working in the field of eukaryotic genome evolution. Since 2023 I am holder of a grant of the Czech Science Foundation (GA ČR) aiming at exploring intron evolution in euglenozoan protists.

Contact:

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