Giulia Magri Ribeiro

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Brazilian/Portuguese, 31 years



Academic CV - September 2025

	Education
2009 2010	ETEC Lauro Gomes, Computer technician
2012 2015	BSc Biological Sciences , Institute of Biosciences - University of São Paulo (hereafter IB-USP), São Paulo, Brazil
2015 2017	Licentiate Biological Sciences, IB-USP, São Paulo, Brazil
2016 2018	MSc in Sciences , <i>IB-USP</i> , São Paulo, Brazil Thesis defense on February 2019
2022	Specialization in Bioinformatics, Unileya, São Paulo, Brazil
<u>2019</u> <u>2023</u>	PhD Zoology , <i>IB-USP</i> , São Paulo, Brazil, Advisor: Dr. Daniel Lahr, Dr. Enrique Lara Thesis defense on August 2023
	Work Experience
	Postdoctoral Researcher in Smith College, Currently working on bioinformatics and single-cell 'omics approaches to study adaptation and population

Postdoctoral Researcher in Bioinformatics, Lund University, Postdoctoral in bioinformatics working with pathogenic bacteria and phenotypic switching. Description of molecular mechanisms of induction and regulation of phenotypic switching in Mycobaterium avium, by using next-generation sequencing approaches and host-pathogen interaction models. Main activities involve: Genome production and Assembly; Transcriptomic analyses by RNA sequencing analysis to compare the gene expression between different stages; Genomic and Methylation analyses focusing on epigenetic regulations, DNA rearrangements that characterize the different stages; HPC computing, Bash, Python and R; Participation and presentation at national and international conferences; Preparation of scientific articles and reports.

genetics of protists. My current research includes next-generation sequencing, genomics, gene expression analysis and confocal microscopy. The goal of my research now is to characterize, life-cycle stages, cryptic species and population

structure of protist lineages living in US bogs and fens environments.

Researcher, Universidade de São Paulo, Exploration of the microbial diversity, through large-scale environmental DNA sequencing techniques (metabarcoding); Laboratory experiments with cell cultures and messenger RNA sequencing (transcriptomes); Evaluation of gene expression by next-generation sequencing and real-time PCR. Bioinformatical analysis of Big data using Python and R. Additionally, activities include participation and presentation at national and international conferences; Teaching and organization of courses; Participation in administrative activities of the university; Preparation of scientific articles and reports.

Visiting scholar, Real Jardín Botánico, CSIC, Madrid, Exploration of microbial diversity in eutrophic environments, through large-scale environmental DNA sequencing techniques (metabarcoding); Laboratory experiments with cell cultures and messenger RNA sequencing (transcriptomes). Bioinformatical analysis using Python and R. Preparation of scientific articles and reports.

2017

Visiting scholar, Smith College, Massachussets, Bioinformatics, processing and analysis of next-generation sequencing data, gene identification, phylogenetic analysis.

2017

Laboratory Technician, Isolation and cultivation of testate amoebae, DNA and RNA extraction and sample preparation for next-generation sequencing

2012 2015

Intern, Universidade de São Paulo, Isolation and cultivation of testate amoebae, DNA extraction, gene amplification, plasmid cloning and Sanger sequencing; Participation and presentation at national and international conferences; Preparation of articles and scientific reports.

Publication List

Laschanzky et al., Laschanzky, K.; Ribeiro, G.M.; Sequeira, R.; Cancade, S.; Carlsson, F.; Lienard, 2025 J. (submitted). Cell wall remodeling-dependent morphotype switch in Mycobacterium avium differentially regulates lung colonization and tissue persistence. Submitted to Nature Microbiology.

Barzilay et al., 2024 Barzilay, D.; Alcino, J.P.; Ribeiro, G.M.; Sousa, A.L.; Lahr, D.J. (2024). Re-evaluating evidence for giant genomes in amoebae. Genetics and Molecular Biology, 47(Suppl 1), e20240092.

Porfirio-Sousa et al., Porfírio-Sousa, A.L; Tice, A.K; Morais, L.; Ribeiro, G. M.; Blandenier, Q.; Dumack. K; Eglit, Y; Fry, N.W; Gomes E Souza, M. B.; Henderson, T.C.; Kleitz-Singleton, F.; Singer D.; Brown, M.W.; Lahr, D.J.G (2024). Amoebozoan testate amoebae illuminate the diversity of heterotrophs and the complexity of ecosystems throughout geological time. Proceedings of the National Academy of Sciences

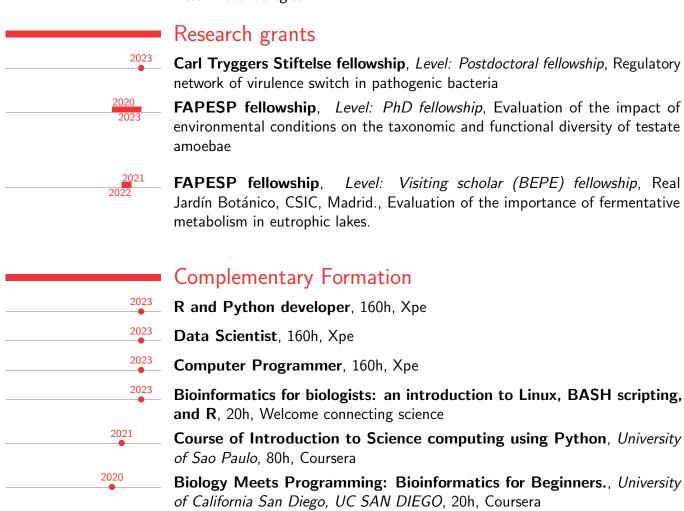
Ribeiro & Lahr, 2024 Ribeiro, G.M.; Lahr, D.J. (2024). Survival in a Changing World: The role of transcriptomics and the urgent need for genomes to understand Arcellinida's adaptive capabilities. Acta Protozoologica, 2024(Special Issue/Early View).

- Ribeiro et al., 2023 Ribeiro, G. M.; Useros, F.; Dumack K.; González-Miguéns R.; Siemensma F.; Porfírio-Sousa, A.F; Soler-Zamora, C.; Alcino J.P.B.; Lahr, D. J. G, Lara E. (2023). Expansion of the cytochrome C oxidase subunit I database and description of four new lobose testate amoebae species (Amoebozoa; Arcellinida) European Journal of Protistology
- Soares et al., 2023 Soares, K. D. A.; Brandão, I; Pereira, J; Gomyde, E; Pessoa-Silva, M; H.; Ribeiro, G. M.; Zanini, F; Grossel, L.A (2023). A decade of Zoology Summer Course: impressions and impacts of the first university extension course on Zoology in Brazil Biota Neotropica
- Ribeiro et al., 2022 Ribeiro, G. M. and Lahr, D. J. G.; (2022) A comparative study indicates vertical inheritance and horizontal gene transfer of arsenic resistance-related genes in eukaryotes. *Molecular Phylogenetics and Evolution*.
- Gonzalez-Miguéns et González-Miguéns, R.; Todorov, M.; Blandenier, Q.; Duckert, C.; Porfirio-Sousa, al., 2022 A. L.; Ribeiro, G. M.; Ramos, D.; Lahr, D. J.G.; Buckley, D.; Lara, E. (2022). Deconstructing Difflugia: The tangled evolution of lobose testate amoebae shells (Amoebozoa: Arcellinida) illustrates the importance of convergent evolution in protist phylogeny Molecular Phylogenetics and Evolution
 - Teng et al., 2021 Teng Y.; Porfírio-Sousa A.L.; Ribeiro, G. M.; Arend M.C.; Meirelles L., Chen E.S., Rosa D.S., Han S.W. (2021). Analyses of the pericyte transcriptome in ischemic skeletal muscles Stem Cell & Research Therapy
 - Hofstatter et al., Hofstatter P. G; Ribeiro, G. M.; Porfírio-Sousa, A. L.; Lahr, D. J. G. (2020)

 2020 The Sexual Ancestor of all Eukaryotes: A Defense of the "Meiosis Toolkit" A Rigorous Survey Supports the Obligate Link between Meiosis Machinery and Sexual Recombination. *BioEssays*.
 - Ribeiro et al., 2019 Ribeiro, G. M.; Porfírio-Sousa, A. L.; Maurer-Alcalá, X. X.; Katz, L. A; Lahr, D. J. G. (2019) De novo sequencing, assembly and annotation of the transcriptome of the free-living testate amoeba Arcella intermedia Journal of Eukaryotic Microbiology.
 - Lahr et al., 2019 Lahr, D. J. G.; Kosakyan, A.; Lara, H.; Mitchell, E. A. D.; Morais, L.; Porfírio-Sousa, A. L.; Ribeiro, G. M.; Tice, A. K.; Kang, S.; Brown, M. W. (2019) Phylogenomics and ancestral morphological reconstruction of testate amoebae demonstrate high diversity of microbial eukaryotes in the Neoproterozoic. Current Biology.
 - Ribeiro et al., 2019 Ribeiro, G. M.; Prado, P.; Coutinho, R.; Rillo, M.; Junior, S.; Porfírio-Sousa, A. L.; Lahr, D. J. G. (2019) Growth rate modulation enables coexistence in a competitive exclusion scenario between microbial eukaryotes. *Acta Protozoologica*.
 - Porfírio-Sousa and Porfírio-Sousa, A. L.; Ribeiro, G. M. and Lahr, D. J. G. (2017). Morphome-tric and genetic analysis of Arcella intermedia and Arcella intermedia laevis (Amoebozoa, Arcellinida) illuminate phenotypic plasticity in microbial eukaryotes. European journal of Protistology

Lahr et al., 2017 Lahr, D. J. G.; Lara, H.; Hofstatter, P. G.; Ribeiro, G. M.; Porfírio-Sousa, A. L.; and Junior, S. P. (2017). Meeting Report: 8th International Symposium on Testate Amoebae, Ilhabela, São Paulo, Brazil, 12–14 September 2016. Journal of Eukaryotic Microbiology

Feres et al., 2016 Feres, J. C.; Porfírio-Sousa, A. L.; Ribeiro, G. M.; Rocha, G. M.; Sterza, J. M.; Souza, M. B. G.; Soares, C. E. A.; and Lahr, D. J. G. (2016). Morphological and morphometric description of a novel shelled amoeba Arcella gandalfi sp. nov.(Amoebozoa: Arcellinida) from Brazilian continental waters. Acta Protozoologica





University of Sao Paulo, 4h, Coursera

Basic principles of genome assembly, annotation & phylogenomics,

Language skills

Portuguese native

2019

English C1

Spanish B2

Academic Quantitative Indicators

Peer-revied Journal 15

articles:

Google Scholar 203, H=8

Citations:

Links

ORCID https://orcid.org/0000-0003-3366-3735

Google Scholar https://scholar.google.com.br/citations?user=3GnK__QAAAAJ&hl=

pt-BR

Lattes http://lattes.cnpq.br/5111122800344440

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Website: https://giuliamagriribeiro.wordpress.com/