Phd, Post doctoral Bioinformatics - Evolutionary Biology & Ecology University of Brussels souza.nataliaa@gmail @nat2bee http://natevolution.com ORCID 0000-0002-0074-6844

Natalia de Souza Araujo

EDUCATION

2012-2017 Ph.D. in Genetics and Evolutionary Biology - University of São Paulo

Expression of Genes Involved in Social Behaviour in Bees with Different Levels of Eusociality. Advisor: Maria Cristina Arias

2010-2012 M.Sc. in Genetics and Evolutionary Biology - University of São Paulo

Analyses of the *Anastrepha fraterculus* complex (Diptera: Tephritidae) in Brazil based on mitochondrial cytochrome oxidase I sequences. Advisor: Andre Luiz Paranhos Perondini

2006-2010 B.Sc. in Biological Science - Universidade Paulista

2004-2007 Technician in Chemistry - Escola Técnica Estadual Getúlio Vargas

LANGUAGE and PROGRAMMING SKILLS

- Portuguese (native); English (fluent); French (intermediary); Spanish (basic)
- Python; R and Unix Environment

RESEARCH EXPERIENCE

- **2019-current** Bioinformatics // Evolutionary Biology & Ecology ULB, University of Brussels (research supervisors: Dr. Serge Aron and Dr. Matthiew Defrance)
- **2017-2019** Unit of Animal Genomics GIGA, University of Liège (research supervisors: Dr. Michel Georges and Dr. Carole Charlier)
- **2012-2017** Laboratory of Genetics and Evolution of Bees, University of São Paulo (research advisor: Dr. Maria Cristina Arias)
- **2014-2015** Laboratory of Ants, evolution & genomics, Queen Mary University of London (research advisor: Dr. Yannick Wurm)
- **2010-2012** Laboratory of Evolution and Genetics of True Fruit Flies, University of São Paulo (research advisor: Dr. Andre Luiz Paranhos Perondini)
- **2008-2010** Laboratory of Evolution and Histophysiology, University of São Paulo (research advisor: Dr. João Carlos Shimada Borges UNIP)

<u>ADVISING</u>

- **2018-current** Paulo Cseri Ricardo Expression of Genes Related to Parasitic Behaviour in Bees. *Coadvisor in PhD Research*
- **2015-2018** <u>Larissa Logullo Piconi</u> Gene Expression Analyses of Behavioural Candidate Genes in Native Bees. *Co-advisor in undergraduate Research*

FUNDINGS

2019 PACBIO, SMRT Leiden – Travel Grant [International]

2016 Society for the Study of Evolution – Travel Grant [International]

2013-2017 FAPESP - Regular Ph.D. Fellowship [National]

2014-2015 FAPESP - BEPE Ph.D. Fellowship Abroad [National]

2010-2012 CNPQ – Regular M.Sc. Fellowship [National]

2008-2010 CNPQ/ UNIP – PIBIC Fellowship for undergraduate students [National]

2006-2010 PROUNI – Scholarship for graduation costs [National]

- **2016** ICE 2nd Place for Best Student Poster. Session: Genetics and Evolutionary Entomology
- **2016** Brazilian Congress of Genetics Honourable Mention for Participation in the Francisco Mauro Salzano Graduate Student Award of Evolution
- **2014** IUSSI 3rd Place for Best Student Poster
- **2011** 57° Brazilian Congress of Genetics Honourable Mention for Participation in the Graduate Student Oral Award of Animal Genetics
- **2009** Instituto Biológico Scientific Merit for Oral Presentation

SELECTED TALKS

- <u>Araujo, N.S.</u>; Wurm Y.; Schmitz B.; Arias M.C. (2018) Unveiling the expression dynamics of genes involved in bee sociality. IUSSI Symposium invited speaker
- <u>Araujo, N.S.</u> and Arias, M. C. (2017) DNA methylation and the evolution of bee eusociality. Evolution 2017
- Araujo, N.S.; Wurm Y.; Arias M.C. (2016) Worker Subcastes: What makes bees nurses? IUSSI-NAS Colloquium

LIST OF PUBLICATIONS

2020

1. **Araujo, N.S.***; Françoso, E.*; Ricardo, P.C.; Santos, P.K.F.; Zuntini, A.R.; Arias, M.C. (2020) Evolutionary perspectives on bee mtDNA from mito-OMICS analyses of a solitary species. Apidologie. doi:10.1007/s13592-020-00740-x

2019

2. **Araujo, N.S.** and Arias M.C. (2019) Mitochondrial genome characterization of *Melipona bicolor*: Insights from the control region and gene expression data. Gene. doi:10.1016/j.gene.2019.04.042

2018

- 3. Santos, P. K. F., **Araujo, N.S.,** Françoso, E., Zuntini, A. R., & Arias, M. C. (2018). Diapause in a tropical oil-collecting bee: molecular basis unveiled by RNA-Seq. BMC genomics, 19(1). doi:10.1186/s12864-018-4694-x
- 4. **Araujo, N. S.**, Santos, P. K. F., & Arias, M. C. (2018). RNA-Seq reveals that mitochondrial genes and long non-coding RNAs may play important roles in the bivoltine generations of the non-social Neotropical bee Tetrapedia diversipes. Apidologie, 49(1), 3-12. doi:10.1007/s13592-017-0542-2

2017

5. **Araujo, N.S.** (2017) Expression of genes involved in the social behaviour of bees with different levels of eusociality. [PhD thesis]. São Paulo: University of São Paulo, Instituto de Biociências. doi:10.11606/T.41.2017.tde-04102017-081728

2016

- 6. **Araujo, N. S.,** Zuntini, A. R., & Arias, M. C. (2016). Getting Useful Information from RNA-Seq Contaminants: A Case of Study in the Oil-Collecting Bee *Tetrapedia diversipes*. Omics, 20(8), 491-492. doi:10.1089/omi.2016.0054
- 7. Arias, M.C. *et al.* (2016) Microsatellite records for volume 8, issue 1. Conservation Genetics Resources. 1(8), 43-81. doi:10.1007/s12686-016-0522-2

2015

8. <u>Araujo, N.S.</u> and Borges J.C.S. (2015) Rodlet cells changes in *Oreochromis niloticus* in response to organophosphate pesticide and their relevance as stress biomarker in teleost fishes. International Journal of Aquatic Biology.

2012

9. <u>Araujo, N.S.</u> (2012) Analyses of the *Anastrepha fraterculus* complex (Diptera: Tephritidae) in Brazil based on mitochondrial *cytochrome oxidase I* sequences. [Masters dissertation]. São Paulo: University of São Paulo, Instituto de Biociências. doi: 10.11606/D.41.2012.tde-18122012-225903