

# *Curriculum Vitae*

## Daniel S. Caetano

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### Appointments

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2022 – present	Towson University, Towson, MD Assistant Professor
2019 – 2021	Universidade de São Paulo, São Paulo, Brazil Postdoctoral researcher Supervisor: Dr. Tiago B. Quental
2017 – 2019	University of Arkansas, Fayetteville, AR Postdoctoral researcher Supervisor: Dr. Jeremy M. Beaulieu

### Education

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2012 – 2017	University of Idaho, Moscow, ID Ph. D. in Bioinformatics and Computational Biology Advisor: Dr. Luke J. Harmon
2009 – 2011	University of São Paulo, São Paulo, Brazil M. S. in Ecology Advisor: Dr. Glauco Machado
2004 – 2009	University of São Paulo, São Paulo, Brazil B. S. in Biological Sciences (major/minor system does not apply)

### Peer reviewed publications

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- Tejero-Cicuéndez H., Patton A.H., **Caetano D.S.**, Šmíd J., Harmon L.J., and S. Carranza, 2022. “Reconstructing Squamate Biogeography in Afro-Arabia Reveals the Influence of a Complex and Dynamic Geologic Past” *Systematic Biology*, 71(2):261–272
- Caetano D.S.** and J.M. Beaulieu, 2020. “Comparative analyses of phenotypic sequences using phylogenetic trees” *American Naturalist*, 192(2) E-Article
- Caetano D.S.** and L.J. Harmon, 2019. “Estimating correlated rates of trait evolution with uncertainty” *Systematic Biology*, 68(3):412–429.
- Caetano D.S.**, O'Meara B.C. and J.M. Beaulieu, 2018. “Hidden state models improve state-dependent diversification approaches, including biogeographical models” *Evolution*, 72(11):2308–2324.
- Caetano D.S.** and L.J. Harmon, 2017. “ratematrix: a R package for studying evolutionary integration among several traits on phylogenetic trees” *Methods in Ecology and Evolution*, 8(12):1920–1927.
- Genevicius B.C., **Caetano D.S.** and C.F. Schwertner, 2017. “Rapid differentiation and asynchronous coevolution of male and female genitalia in stink bugs” *Journal of Evolutionary Biology*, 30(3):461–473.
- Uyeda J.C., **Caetano D.S.** and M.W. Pennell, 2015. “Comparative analysis of Principal Components can be misleading” *Systematic Biology*, 64(4):677–689.

- Caetano D.S.** and A. Aisenberg, 2014. "Forgotten treasures: the fate of data in animal behaviour studies" *Animal Behaviour*, 98:1–5.
- Caetano D.S.** and G. Machado, 2013. "The ecological tale of Gonyleptidae (Arachnida, Opiliones) evolution: phylogeny of a Neotropical lineage of armoured harvestmen using ecological, behavioural and chemical characters" *Cladistics*, 29(6):589–609.
- Caetano D.S.**, Bená D.D.C. and S.A. Vanin, 2013. "*Copelatus cessaima* sp. nov. (Coleoptera: Dytiscidae: Copelatinae): first record of a troglomorphic diving beetle from Brazil" *Zootaxa*, 3710(3):226–232.
- Werneck R.M., **Caetano D.S.** and G. Machado, 2012. "Maternal care in the Neotropical harvestman *Liogonyleptoides tetracanthus* (Opiliones: Gonyleptidae)" *Journal of Arachnology*, 40(1):135–137.
- Caetano D.S.** and S.A. Vanin, 2008. "A new species of the genus *Pimelerodius* Vanin, 1986 from the Amazon Region, with notes on the geographic distribution of *Pimelerodius motacilla* (Boheman, 1843) (Coleoptera, Curculionidae, Otidoccephalini)" *Revista Brasileira de Entomologia*. 52(3):431–433.

### Manuscripts in preparation

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- Caetano D.S.**, Uyeda J.C., Harmon L.J., and R. Zenil-Ferguson. "Using predictive functions as continuous regimes for rates of trait evolution across a phylogeny" *In preparation for Systematic Biology*.
- Caetano D.S.** and T.B. Quental. "How important is budding speciation for phylogenetic comparative models?" Resubmitted to *Systematic Biology*. Pre-print available here: <https://doi.org/10.1101/2022.05.24.493296>

### Software

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- 'CorrSeq' (Open source) – <https://github.com/Caetanods/CorrSeq>  
*R package for fitting phylogenetic comparative models of trait evolution for sequence phenotypes.*
- 'hisse' (Open source) – <https://cran.r-project.org/web/packages/hisse/index.html>  
*R package for testing models of trait-dependent diversification using Hidden Markov models.*
- 'ratematrix' (Open source) – <https://cran.r-project.org/web/packages/ratematrix/index.html>  
*R package for the study of evolutionary integration using phylogenies in a Bayesian framework.*
- 'Posterior\_Phylo\_sampler' (Open source) – [https://github.com/Caetanods/Posterior\\_Phylo\\_sampler](https://github.com/Caetanods/Posterior_Phylo_sampler)  
*A memory efficient bash script to randomly sample phylogenies from large nexus files.*
- Programming languages: R (advanced); C++ (intermediary); python (intermediary); bash (intermediary)

### Outreach & Digital media

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- Caetano D. S.** 2017. "Phylogenies, trait evolution and fancy glasses" Invited blog post on *Methods in Ecology and Evolution* - <https://methodsblog.wordpress.com/2017/10/19/phylogenies-trait-evolution/>
- Twitter account (@ds\_caetano)**: I share and discuss research with 1,024 followers.
- Recorded talk on YouTube**: (Evolution 2019) <https://www.youtube.com/watch?v=9T3XF5Fq7xE>

### Awards and fellowships

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| 2023        | Society of Systematic Biologists Symposium award - \$10,000                                   |
| 2019        | Top-reviewer for <i>Evolution</i> (SSE).  |
| 2016 - 2017 | Bioinformatics and Computational Biology Program fellowship - \$48,050<br>University of Idaho |

- 2012 - 2016 Science Without Borders Fellowship - \$157,560  
CAPES, Brazil
- 2015 College of Sciences Travel Award - \$1,300  
University of Idaho
- 2012 - 2013 GPSA Travel Award - \$1,400  
University of Idaho
- 2009 - 2011 FAPESP Master level Scholarship - \$14,400  
São Paulo state, Brazil
- 2008 – 2009 FAPESP Undergraduate level Scholarship - \$2,316  
São Paulo state, Brazil

### **Invited talks**

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- 2023 *Systematics Symposium* “Understanding the Drivers of (Insect) Diversity through the Integration of Phylogenies and Natural History Data”  
Eastern Branch Meeting of the Entomological Society of America (Providence, RI)  
Title: “Using phylogenies to understand phenotypic evolution in insects”
- 2022 University of Maryland  
College Park, MD  
Title: “Phylogenetic comparative methods and the macroevolution of complex phenotypes”
- 2022 University of Maryland, Baltimore County  
Baltimore, MD  
Topic: Discussion with students about the academia job market, net-working, and preparation for interviews
- 2018 Jetz lab – Research Updates  
Yale University, New Haven, CT  
Title: “Hidden state models and rate variation on phylogenetic trees”
- 2016 External IBEST Evaluation – Research Updates  
University of Idaho, Moscow, ID  
Title: “Comparative methods to study the evolution of correlated traits using phylogenetic trees”
- 2014 and 2015 Biology 489 (Herpetology)  
University of Idaho, Moscow, ID  
Title: "Evolution of warning signals and mimicry in Neotropical snakes"
- 2011 Neotropical Harvestmen symposium  
III Latin-American Congress of Arachnology, Montenegro, Colombia  
Title: “The ecological tale of Gonyleptidae evolution: phylogeny using behavioral, ecological, and chemical characters”
- 2011 Seminários da Ecologia (Graduate seminar series)  
University of São Paulo, São Paulo, Brazil  
Title: “Binary animal behavior: phylogeny of a harvestmen family and evolutionary patterns of behavioral characters”
- 2011 Biological systematics class  
Federal University of São Paulo, São Paulo, Brazil  
Title: “Use of behavioral characters in phylogenetic inference”

### **Conference presentations**

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- 2019 Evolution Meeting  
Title: “Comparative analyses of phenotypic sequences: a case study using cricket songs”
- 2018 Botany Meeting  
Title: “A new hope: hidden state models improve the adequacy of state-dependent diversification approaches using empirical trees”

- 2017 Society of Systematic Biologists – Standalone Meeting  
Title: “ratematrix: a R package for studying evolutionary integration among several traits on phylogenetic trees” (lightning talk)
- 2016 Evolution Meeting  
Title: “Estimating correlated rates of trait evolution from phylogenies with uncertainty”
- 2015 Ernst Mayr symposium (Evolution Meeting - SSB Student Award competition)  
Title: “The colors of deception: Evolution of warning signals in Neotropical snakes (Colubroidea: Dipsadidae)”

## **Teaching and student supervision**

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### *Semester-long courses*

- since 2023 BIOL208 – Biodiversity (Towson University)  
since 2022 MBBS 301 – Introduction to Bioinformatics (Towson University)  
since 2022 BIOL206L – Introduction to Ecology and Evolution (Towson University)

### *Short courses and workshops*

- 2021 Short course – Introduction to statistical modeling in R (Universidade de São Paulo)
- 2020 Workshop instructor, Cleveland, OH (*Evolution Meeting 2020*)  
“Using Simulations in Phylogenetics and Comparative Studies”
- 2020 Workshop instructor, Gainesville, FL (Meeting of the *Society of Systematic Biologists*)  
“State-dependent diversification models via graphical models and RevBayes”
- 2017 Workshop instructor, University of São Paulo  
“MCMC step by step: Applications of Monte Carlo methods in phylogenetic comparative methods”
- 2016 Workshop instructor, University of São Paulo  
“MCMC step by step: Applications of Monte Carlo methods in phylogenetic comparative methods”

### *Teaching assistantships*

- 2018 Assistant instructor, University of Arkansas  
“Molecular Phylogenetics” (undergraduate / graduate level)
- 2015 Workshop helper, Washington State University, Pullman, WA  
“Software carpentry”
- 2014 Teaching assistant, University of Idaho  
“Computer Skills for Biologists”
- 2011 Teaching assistant, University of São Paulo  
“Use of the R Language to analyze ecological data”
- 2010 Teaching assistant, University of São Paulo  
“Introduction to Ecology” (undergraduate level)

### *Student supervision*

- 2018 - 2020 Supervision of M. S. student Peter Hasik at the University of Arkansas  
2018 - 2020 Supervision of Ph. D. student James Boyko at the University of Arkansas

### *Student mentorship*

- since 2022 Mentoring 11 undergraduate students (Towson University)  
since 2023 Selected mentor for graduate students by the *Society of Systematic Biologists*.

## Reviewer

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I am a reviewer for *Systematic Biology*; *Proceedings of the Royal Society B: Biological Sciences*; *The American Naturalist*; *Ecology Letters*; *Evolution*; *Methods in Ecology and Evolution*; *PLOS One*; *PLOS Computational Biology*; *Evolutionary Bioinformatics*; *American Journal of Botany*; *Biological Journal of the Linnean Society*; *Nature Communications*; *Zoologia*; *Revista Brasileira de Entomologia*; *Austral Ecology*; *BMC Evolutionary Biology*; *Journal of Insect Behavior*; *Journal of Natural History*; *New Phytologist*; *PeerJ*

I also review grant submissions for CONICYT (Chile).

## Professional service

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2023 -	Committee member of the Student Awards and Scholarships Towson University, Towson, MD
2015 - 2017	Committee member of the Bioinformatics and Computational Biology seminar series University of Idaho, Moscow, ID
2015 - 2016	Graduate student representative Bioinformatics and Computational Biology program (University of Idaho)
2011	R study group – Introduction to the R language (Graduate level) University of São Paulo, São Paulo, Brazil
2009 - 2010	R study group – Phylogenies and the R language (Graduate level) University of São Paulo, São Paulo, Brazil
2010 - 2011	Committee member of the Department of Ecology seminars University of São Paulo

## Synergistic activities

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2023	Organizer of the Society of Systematic Biologists Symposium at the 2023 <i>Evolution</i> meeting (Albuquerque, NM). Title: “Paving the road to the phenome: challenges in working with a large number of phenotypes”
2018	ARBOR Hackaton: Cornell University Organizers: Dr. Chelsea Specht and Dr. Luke Harmon
2016 and 2017	Behavior Data Meeting; Cornell Lab of Ornithology Organizer: Dr. Mike Webster

## Consulting service

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2011-2012	Project manager – Collection and identification of insects (Coleoptera) Location: Cerrado region (Belo Horizonte, Brazil)
2011-2012	Field assistant – Collection of insects (Lepidoptera) Location: Amazon region (Rondônia, Brazil)