

# Basanta Khakurel, M.S.

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### **Education**

2023 – Present

Ph.D., Ludwig Maximilians-Universität München.

Advisor: Dr. Sebastian Höhna

Thesis topic: Robust models for morphological phylogenetic inferences.

2021 - 2023

M.S. Biology, Southeastern Louisiana University.

Advisor: Dr. April Wright

Thesis title: Inference of Phylogeny using Discrete Morphological Characters.

2017 - 2021

B.S. Biology, Southeastern Louisiana University.

Minors: Chemistry and Political Science

Advisor: Dr. April Wright

Thesis title: Species delimitation of Eastern Pinesnakes (Pituophis melanoleucus).

# **Employment History**

2023 – Present Research Assistant, Department of Earth and Environmental Sciences, Ludwig-Maximilians-Universität München.

2019 – 2021 Undergraduate Research Assistant, Department of Biological Sciences, Southeastern Louisiana University.

2019 – 2019 Summer Camp Teacher, Upward Bound Math and Science Program, Southeastern Louisiana University.

2018 – 2019 **Student Assistant,** Sims Memorial Library, Southeastern Louisiana University.

#### **Presentations and Talks**

#### **Talks**

- **B. Khakurel**, "Snakes, fish, and bayesian morphological phylogenetics," in *PhD and PostDoc Seminar*, Paleontological Museum Munich, 2024.
- **B. Khakurel**, B. M. Wynd, and A. M. Wright, "Use of continuous traits in phylogenetic estimation," in *Departmental Seminar*, Southeastern Louisiana University, 2023.
- **B. Khakurel**, "Simulations in biology," in *Biological Data Analysis Class*, Southeastern Louisiana University, 2022.

#### **Presentations at Academic Conferences**

- **B. Khakurel** and S. Höhna, "Characters changing speed: The covariomorph model and its impact on phylogenetic trees," in *Evolution Meetings*, Athens, Georgia, USA, 2025.
- **B. Khakurel** and S. Höhna, "An efficient heterotachy model for morphological phylogenetics," in 3rd Joint Congress on Evolutionary Biology, Montréal, Canada, 2024.
- **B. Khakurel** and S. Höhna, "Evidence of covarion rate variation in empirical morphological datasets," in *The Paleontological Association Annual Meeting*, Poster presentation, Erlangen, Germany, 2024.

- N. Heckeberg, A. Capobianco, G. Darlim, **B. Khakurel**, and S. Höhna, "Potential and perils in using morphological characters for phylogenetic inference," in *The Palaeontological Association Annual Meeting*, Poster presentation, Cambridge, UK, 2023.
- **B. Khakurel** and A. M. Wright, "Model adequacy in morphological data," in *Society of Systematic Biologists Meeting*, Mexico City, Mexico, 2023.
- **B. Khakurel**, B. M. Wynd, and A. M. Wright, "Use of continuous traits in phylogenetic estimation," in *Evolution Meetings*, Albuquerque, NM, USA, 2023.
- B. M. Wynd, **B. Khakurel**, and A. M. Wright, "Incorporating continuous characters in total-evidence phylogenetic estimation of extinct lineages," in *GSA Connects Meeting*, Pittsburgh, PA, USA, 2023.
- **B. Khakurel** and A. M. Wright, "Species delimitation in eastern pine snakes (*Pituophis melanoleucus*)," in *Evolution Meetings*, Cleveland, OH, USA, 2022.
- 9 O. Schwery, **B. Khakurel**, and A. M. Wright, "Adequacy of fossilised birth-death models: A test case in ants," in *Evolution Meetings*, Cleveland, OH, USA, 2022.
- **B. Khakurel** and A. M. Wright, "Species delimitation in eastern pine snakes (*Pituophis melanoleucus*)," in *LBRN 8th Computational Biology and Bioinformatics Meeting*, Baton Rouge, LA, USA, 2021.
- **B. Khakurel**, C. Grigsby, T. Tran, and A. M. Wright, "Character state space partitioning in phylogenetic analyses from discrete data," in *Louisiana Biomedical Research Network 18th Annual Meeting*, Poster presentation, Baton Rouge, LA, USA, 2020.
- **B. Khakurel**, C. Grigsby, and A. M. Wright, "Modeling morphological state space," in *Society of Systematic Biologists Meeting*, Poster presentation, Gainesville, FL, USA, 2020.
- **B. Khakurel**, V. Rocquin, and J. Litolff, "The role of honors student organizations in building and sustaining a robust honors presence on campus," in *Southern Regional Honors Council Conference*, Memphis, TN, USA, 2019.

#### Research Publications

## **Submitted Preprints**

- **B. Khakurel** and S. Höhna, A covarion model for phylogenetic estimation using discrete morphological datasets, 2025.
- B. M. Wynd, **B. Khakurel**, C. F. Kammerer, P. J. Wagner, and A. M. Wright, *Incorporating continuous characters in joint estimation of dicynodont phylogeny*, 2025.

#### **Journal Articles**

- N. S. Heckeberg, A. Capobianco, **B. Khakurel**, G. Darlim, and S. Höhna, "Practical guide and review of fossil tip-dating in phylogenetics," *Systematic Biology*, 2025 (in press).
- J. C. G. Duque, M. Balk, W. Dahdul, *et al.*, "Meeting report for the phenoscape traitfest 2023 with comments on organising interdisciplinary meetings," *Biodiversity Information Science and Standards*, vol. 8, e115232, 2024. ODI: https://doi.org/10.3897/biss.8.115232.
- **B. Khakurel**, C. Grigsby, T. D. Tran, J. Zariwala, S. Höhna, and A. M. Wright, "The fundamental role of character coding in bayesian morphological phylogenetics," *Systematic biology*, vol. 73, no. 5, pp. 861–871, 2024. ODI: https://doi.org/10.1093/sysbio/syae033.
- **B. Khakurel**, Z. L. Nikolakis, B. I. Crother, and A. Wright, "Species delimitation of eastern pinesnake complex (pituophis melanoleucus)," *Bulletin of the Society of Systematic Biologists*, vol. 2, no. 1, 2023.

  \*\*DOI: https://doi.org/10.18061/bssb.v2i1.9423.

L. M. Ward, C. D. McMahan, **B. Khakurel**, A. M. Wright, and K. R. Piller, "Genomic data support the taxonomic validity of middle american livebearers poeciliopsis gracilis and poeciliopsis pleurospilus (cyprinodontiformes: Poeciliidae)," *PloS one*, vol. 17, no. 1, e0262687, 2022. ODOI: https://doi.org/10.1371/journal.pone.0262687.

#### Skills

Languages Strong reading, writing and speaking competencies for Nepali, English, and Hindi. Basic German understanding.

Coding Python, R, LTFX, C++

Misc. Academic research, teaching, training, Large typesetting and publishing.

# Miscellaneous Experience

### **Professional Memberships**

- Society for the Study of Evolution (SSE)
- Society of Systematic Biologists (SSB)
- Phi Kappa Phi Honor Society

#### Awards, Fellowships, and Grants

- 2023 SSB Travel Award, Society of Systematic Biologists.
- Distinguished Graduate Fellowship, Department of Biological Sciences and the Dean of the College of Science and Technology, Southeastern Louisiana University.
  - **Graduate Research Fellowship**, Louisiana Biomedical Research Network.
- David Ware Outstanding International Student Award, Southeastern Louisiana University.
  - Outstanding Senior Award, Department of Biological Sciences, Southeastern Louisiana University.
  - **Dr. Marvin L. Yates Award of Excellence**, Southeastern Louisiana University.
- **Sophomore Honors Distinction**, Honors Department, Southeastern Louisiana University.

#### Service and Outreach

2018 - 2022 President, Nepalese Student Association at Southeastern

Planning Committee, Louisiana Sci-Comm Summit

2020 - 2021 Vice President, Phi Kappa Phi

2019 - 2021 Ambassador, Honors Program, Southeastern Louisiana University

2019 - 2020 Leadership Ambassador, Division of Student Affairs

2019 Senator, Student Government Association

2018 Volunteer Judge, Science Fair at Southeastern Lab School

### **Professional Development**

Navigating Scientific Publishing – Early Career Focus, workshop was offered in the 3rd Joint Congress on Evolutionary Biology.

In this workshop, the organizers helped early career researchers understand the publish-

ing landscape and how to efficiently navigate it to further your professional life.

# Miscellaneous Experience (continued)

- Phenoscape Traitfest, was a hands on workshop that aimed at methods generating, discovering, linking to, recombining, or computing with machine-interpretable trait data. In this workshop, I learned processes to expedite the process of data acquisition and implement machine-learning approached to automate the collection of measurements on large datasets.
  - RevBayes Workshop and Hackathon, was a workshop/hackathon offered in the Munich on October, 2023.

    In this workshop, I learned about about Bayesian Phylogenetic Methods, how to model phylogenetic analyses in RevBayes and also about the use of C++ for academic software development.
- PhylogatR (Phylogeographic data Aggregation and Repurposing) workshop, was a workshop for the beta-testing of the phylogatR database. It is a database that brings together genetic data with georeferenced specimen records that are analysis-ready. In this workshop, I learned the use of phylogatR database and got a chance to connect with researchers working in various fields.
- Posterior Predictive Simulation with P2C2M workshop, was a workshop offered in the Society of Systematic Biologists meeting 2020 in Gainesville, FL.

  In this workshop, I learned about posterior predictive simulations which is a powerful tool in evaluating model performance in Bayesian Phylogenetic Methods.

### Mentoring

- 2020-2023 Simran Baruwal, Undergraduate Research Assistant, Southeastern Louisiana University
  - Anisha Neupane, Undergraduate Research Assistant, Southeastern Louisiana University

## References

Available on Request