Lena Collienne

Curriculum Vitae



Research

Postdoctoral Research, Fred Hutch Cancer Center.

Developing novel methods for phylogenetic inference using machine learning.

Postdoctoral Research, *University of Canterbury*.

Extending subtree prune and regraft operations to ranked phylogenetic trees and investigating thereby introduced distance measures for phylogenetic time trees.

PhD Research, University of Otago.

Introducing and analysing spaces of phylogenetic time trees based on nearest neighbour interchange tree rearrangements.

M.Sc. Research, University of Greifswald.

Establishing properties of ranked nearest neighbour interchange moves between ranked phylogenetic trees

Education

2016–2018 Master of Science, Biomathematics, University of Greifswald (GER).

2012–2015 Bachelor of Science, Biomathematics, University of Greifswald (GER).

Work Experience

2023-now Postdoctoral Research Fellow, Matsen group, Fred Hutch Cancer Center (US).

2022–2023 **Postdoctoral Research Fellow**, *BioDS lab*, *School of Mathematics and Statistics*, University of Canterbury (NZ).

2022 **Lecturer**, STAT211: Random Processes, University of Canterbury (NZ).

2019 **Tutor**, COSC341: Theory of Computing, University of Otago (NZ).

2015–2016 **Summer Research Project**, *University of Auckland (NZ)*.

Scholarships and Awards

2022 Hatherton Award (Royal Society of New Zealand)

2021 Exceptional PhD thesis (Division of Science, University of Otago)

2018–2021 University of Otago Doctoral Scholarship

2018 Externally Funded Research Grant (Max Planck Institute Plön)

2015 Summer Research Scholarship (University of Auckland)

- 2015 PROMOS Travel Scholarship (University of Greifswald)
- 2014–2015 Deutschlandstipendium (Alfried Krupp von Bohlen und Halbach Foundation/Federal Government of Germany)

Additional Activities

- 2021 President of the Otago Computer Science Society (University of Otgao)
- 2019–2021 Member of the Postgraduate Committee (Department of Computer Science, University of Otago)
- 2019–2021 Organising the annual Postgraduate Symposium (Department of Computer Science, University of Otago)
- 2019–2021 Member of Student Council (Institute for Mathematics and Computer Science, University of Greifswald)

Talks

Conference Talks

- 2024 **Evolution 2023**, *Montreal (CA)*.

 Contributed talk: Instability in Phylogenetic Trees after Taxon Addition
- 2023 **SMB meeting 2023**, *Columbus (OH, US)*. Invited minisymposium talk: Spaces of Discrete Time Trees
- 2022 **Phylomania 2022**, *Hobart (AU)*.

 Contributed talk: Subtree Prune and Regraft on Ranked Trees
- 2021 **Phylomania 2021** (Best Student Talk Award), *Online*. Contributed talk: Distances between Phylogenetic Time Trees
- 2021 **NZ Phylogenomics Meeting**, *Akaroa (NZ)*. Contributed talk: The Space of Discrete Coalescent Trees
- 2020 **NZ Phylogenomics Meeting**, *Waiheke (NZ)*.

 Contributed talk: Online Algorithms in Computational Biology
- 2019 NZ Phylogenomics Meeting, Napier (NZ).
 Contributed talk: The Ranked Nearest Neighbour Interchange Space of Phylogenetic Trees
 Invited Seminar Talks
- 2021 **Departmant of Mathematics**, *University of Otago (NZ)*. The Space of Discrete Coalescent Trees
- 2020 Online Seminars on Algorithms and Complexity in Phylogenetics, Online.
 Computing the Ranked Nearest Neighbour Interchange Distance between Ranked Phylogenetic Trees
- 2019 Max Planck Instityte for Mathematics in the Science, Leipzig (GER). The Ranked Nearest Neighbour Interchange space of phylogenetic trees
- 2017 **Computational Evolution Group**, *ETH Zurich (CH)*. Discrete Time Trees

Other

- 2022 **School of Mathematics and Statistics**, *University of Canterbury (NZ)*. How to Give a (Good) Talk
- 2020 **Postgraduate Symposium** (1st place Best Presentation Award), *University of Otago (NZ)*.
 - The Complexity of Computing the RNNI Distance between Phylogenetic Trees
- 2020 **Seminar of Departments of Computer Science and Information Science**, *University of Otago (NZ)*.
 - The Complexity of Computing Nearest Neighbour Interchange Distances between Ranked Phylogenetic Trees
- 2019 **Postgraduate Symposium** (2nd place Best Presentation Award), *University of Otago (NZ)*.
 - Online Algorithms in Computational Biology

Publications

Collienne, L., Barker, M., Suchard, MA., & Matsen IV, FA. (2024). Phylogenetic tree instability after taxon addition: empirical frequency, predictability, and consequences for online inference. *Systematic Biology syae059*. https://doi.org/10.1093/sysbio/syae059

Collienne, L., Whidden, C. & Gavryushkin, A. (2024). Ranked Subtree Prune and Regraft. Bulleting of Mathematical Biology 86, 24. https://doi.org/10.1007/s11538-023-01244-2

Berling, L., **Collienne, L.** & Gavryushkin, A. (2023). Estimating the mean in the space of ranked phylogenetic trees. *BioRxiv*. https://doi.org/10.1101/2023.05.08.539790.

Bouckaert, R., **Collienne, L.** & Gavryushkin, A. (2022). Online Bayesian Analysis with BEAST2. *BioRxiv*.

Collienne, L. (2021). Spaces of phylogenetic time trees (Thesis, Doctor of Philosophy). University of Otago. Retrieved from http://hdl.handle.net/10523/12606

Collienne, L., Elmes, K., Fischer, M., Bryant, D. & Gavryushkin, A. (2021). Discrete Coalescent Trees. *Journal of Mathematical Biology 83.5, p. 60. issn:* 1432-1416.

Collienne, L. & Gavryushkin, A. (2021). Computing nearest neighbour interchange distances between ranked phylogenetic trees. *Journal of Mathematical Biology 82.1, p. 8. issn: 1432-1416.*