

# CURRICULUM VITAE

1/2024

## PERSONAL DETAILS

Name	Tuomas Hämälä
Nationality	Finland
Email address	tuomas.hamala@luke.fi
Public profiles	<a href="#">Google Scholar</a> , <a href="#">ResearchGate</a>
ORCID iD	<a href="#">0000-0001-8306-3397</a>
Webpage	<a href="#">thamala.github.io</a>

## EMPLOYMENT

1/2024 – present	Natural Resources Institute Finland (Luke), Finland <b>Senior scientist in plant genetics and genomics</b>
4/2022 – 12/2023	University of Nottingham, UK <b>Marie Skłodowska-Curie individual fellow</b> Supervisor: Levi Yant
9/2021 – 3/2022	University of Oulu, Finland <b>Academy of Finland postdoctoral fellow</b>
8/2018 – 7/2021	University of Minnesota Twin Cities, USA <b>Postdoctoral associate</b> Supervisor: Peter Tiffin

## EDUCATION

1/2014 – 7/2018	University of Oulu, Finland <b>PhD</b> in population genetics Thesis title: “Ecological genomics in <i>Arabidopsis lyrata</i> : local adaptation, phenotypic differentiation, and reproductive isolation” Supervisor: Outi Savolainen
9/2011 – 12/2013	University of Oulu, Finland <b>MSc</b> in genetics
9/2008 – 5/2011	University of Oulu, Finland <b>BSc</b> in bioscience

## PUBLICATIONS

\*equal contribution, ✉corresponding author

13. **Hämälä T**✉, Moore C, Cowan L, Carlile M, Gopaulchan D, Brandrud MK, Birkeland S, Loose M, Kolář F, Koch MA & Yant L ✉ (2023). Impact of whole-genome duplications on structural variant evolution in the plant genus *Cochlearia*. *bioRxiv* preprint: 2023/560073.
12. Bray SM, **Hämälä T**, Zhou M, Busoms S, Fischer S, Desjardins SD, Mandáková T, Moore C, Mathers TC, Cowan L, Monnahan P, Koch J, Wolf EM, Lysak MA, Kolář F, Higgins JD, Koch MA & Yant L ✉ (2023). Kinetochore and ionomic adaptation to whole genome duplication. *bioRxiv* preprint: 2023/559727.
11. Epstein B, Burghardt L, Heath K, Grillo M, Kostanecki A, **Hämälä T**, Young N & Tiffin P ✉ (2023). Combining GWAS and population genomic analyses to characterize coevolution in a legume-rhizobia symbiosis. *Molecular Ecology* 23: 3798–3811.
10. **Hämälä T**✉, Ning W, Kuittinen H, Aryamanesh N\* & Savolainen O\* (2022). Environmental response in gene expression and DNA methylation reveals factors influencing the adaptive potential of *Arabidopsis lyrata*. *eLife* 11:e83115.
9. **Hämälä T**\*✉, Wafula EK\*, Guiltinan MJ, Ralph PE, dePamphilis CW & Tiffin P ✉ (2021). Genomic structural variants constrain and facilitate adaptation in natural populations of *Theobroma cacao*, the chocolate tree. *PNAS* 118: e2102914118.
8. Takou M, **Hämälä T**, Koch E, Steige KA, Dittberner H, Yant L, Genete M, Sunyaev S, Castric V, Vekemans X, Savolainen O & de Meaux J ✉ (2021). Maintenance of adaptive dynamics and no detectable load in a range-edge out-crossing plant population. *Molecular Biology and Evolution* 38: 1820–1836.
7. **Hämälä T**✉ & Tiffin P (2020). Biased gene conversion constrains adaptation in *Arabidopsis thaliana*. *Genetics* 215: 831–846.
6. **Hämälä T**\*✉, Gorton AJ\*, Moeller AD & Tiffin P ✉ (2020). Pleiotropy facilitates local adaptation to distant optima in common ragweed (*Ambrosia artemisiifolia*). *PLOS Genetics* 16: e1008707.
5. **Hämälä T**✉, Guiltinan MG, Marden JH, Maximova S, dePamphilis C & Tiffin P ✉ (2020). Gene expression modularity reveals footprints of polygenic adaptation in *Theobroma cacao*. *Molecular Biology and Evolution* 1: 110–123.
4. **Hämälä T**✉ & Savolainen O ✉ (2019). Genomic patterns of local adaptation under gene flow in *Arabidopsis lyrata*. *Molecular Biology and Evolution* 36: 2557–2571.
3. Mattila TM\*, Laenen B\*, Horvath R, **Hämälä T**, Savolainen O & Slotte T ✉ (2019). Impact of demography on linked selection in two outcrossing Brassicaceae species. *Ecology and Evolution* 9: 9532–9545.
2. **Hämälä T**✉, Mattila TM & Savolainen O (2018). Local adaptation and ecological differentiation under selection, migration, and drift in *Arabidopsis lyrata*. *Evolution* 72: 1373–1386.
1. **Hämälä T**✉, Mattila TM, Leinonen PH, Kuittinen H & Savolainen O (2017). Role of seed germination in adaptation and reproductive isolation in *Arabidopsis lyrata*. *Molecular Ecology* 26: 3484–3496.

## FUNDING

- 2021 European Commission, Marie Skłodowska-Curie individual fellowship, 213 000€
- 2021 Academy of Finland, postdoctoral fellowship, 330 000€
- 2019 MPG, travel grant, \$800
- 2018 Emil Aaltonen foundation, travel grant, 3000€
- 2018 SMBE, young investigator travel award, \$2000
- 2017 Emil Aaltonen foundation, research fellowship, 13 350€
- 2017 UniOGS, travel grant, 1500€
- 2015 PGDP, travel grant, 500€

## SKILLS AND EXPERTISE

- Population genetics, ecology, and genomics: local adaptation, polygenic adaptation, structural variants, genome evolution, speciation
- Bioinformatics: > 10 years of experience with high-performance computing and next-generation sequence analysis; familiar with both model (whole-genome, transcriptome, long-reads, assemblies) and non-model (transcriptome, RAD-Seq) organism; extensive experience with individual-based simulations (coalescent and forward-in-time); fluent in programming language C
- Statistics: fluent in R; regression analysis (e.g., linear mixed models, model selection); machine-learning (e.g., random forest, neural networks); data visualization
- Field and laboratory experiments: designed and maintained large-scale reciprocal transplant, common garden, and growth-room experiments

## TEACHING AND MENTORING EXPERIENCE

- 2023 MSc thesis supervision  
Xiuzhiping Quan (bioinformatics major)
- 2022 MSc thesis co-supervision  
Ana Da Silva, Matthew Gaskins (bioinformatics majors)
- 2015 – 2016 MSc thesis supervision  
Margarita Takou (ecology and population genetics major)
- 2013 – 2016 Practical trainee supervision  
Petri Vänni, Weixuan Ning, Elina Haataja, Margarita Takou, Rami-Petteri Apuli, Paul DuBray, Toni Jenfors
- 2015 Co-instructor  
Graduate-level workshop on population genomics, 6 h
- 2014 Teaching assistant  
Basics in genetics, 15 h

## PRESENTATIONS

### Invited departmental seminars and conference talks

8/2023	Department of Ecology & Evolution, University of Sussex, UK
5/2023	Department of Evolution, Ecology & Behaviour, University of Liverpool, UK
3/2023	Milner Centre for Evolution, University of Bath, UK
11/2021	Agricultural Genomics Institute at Shenzhen, online
11/2020	Department of Ecology & Genetics, University of Oulu, online
1/2020	PAG conference, San Diego, California

### Contributed conference talks and posters

7/2023	Oral, SMBE, Ferrara, Italy
8/2022	Oral, ESEB, Prague, Czech Republic
1/2022	Oral, PopGroup, online
8/2019	Poster, ESEB, Turku, Finland
8/2018	Oral, Midwest PopGen meeting, Saint Paul, Minnesota
7/2018	Poster, SMBE, Yokohama, Japan
2/2018	Oral, Arabis symposium, Cologne, Germany
7/2017	Oral, SMBE, Austin, Texas
8/2015	Poster, ESEB, Lausanne, Switzerland

## REFERENCES

- Levi Yant (postdoctoral supervisor)  
Professor at University of Nottingham, UK, [levi.yant@nottingham.ac.uk](mailto:levi.yant@nottingham.ac.uk)
- Peter Tiffin (postdoctoral supervisor)  
Professor at University of Minnesota Twin Cities, USA, [ptiffin@umn.edu](mailto:ptiffin@umn.edu)
- Outi Savolainen (PhD supervisor)  
Emerita professor at University of Oulu, Finland, [outi.savolainen@oulu.fi](mailto:outi.savolainen@oulu.fi)

## OTHER

### Journal referee

Annals of Botany, eLife, Evolution, Evolution Letters, Evolutionary Applications, Evolutionary Ecology, Frontiers in Plant Science, G3: Genes | Genomes | Genetics, Journal of Heredity, Molecular Biology and Evolution, Molecular Ecology, Molecular Ecology Resources, Molecular Plant, New Phytologist, The Plant Cell, PLOS Genetics, PNAS

Programs and scripts written for NGS data analysis: [GitHub](#)