

CURRICULUM VITAE  
6/2021

PERSONAL DETAILS

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

EMPLOYMENT

8/2018 – present	University of Minnesota Twin Cities, USA <b>Postdoctoral associate</b> Supervisor: Peter Tiffin
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EDUCATION

1/2014 – 7/2018	University of Oulu, Finland <b>PhD</b> in population and evolutionary 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

\*Corresponding author, †Equal contribution

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 (in press).
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\***<sup>†</sup>, Gorton AJ<sup>†</sup>, 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<sup>†</sup>, Laenen B<sup>†</sup>, 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€ (accepted in part)
- 2019 MPG, travel grant – \$800
- 2018 Emil Aaltonen foundation, travel grant – 3000€
- 2018 SBE, young investigator travel award – \$2000
- 2017 Emil Aaltonen foundation, research fellowship – 13,350€
- 2017 UniOGS, travel grant – 1500€
- 2015 Population genetics doctoral program, travel grant – 500€

## SKILLS AND EXPERTISE

Population genetics: local adaptation, polygenic adaptation, structural variants, genome evolution, speciation

Bioinformatics: >8 years of experience with high-performance computing and next-generation sequence analysis; familiar with both model (whole-genome and transcriptome) and non-model (transcriptome and 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

- 2015 – 2016 Master's thesis supervision  
Margarita Takou, ecology and population genetics
- 2013 – 2016 Practical trainee supervision  
Petri Vänni, Weixuan Ning, Elina Haataja, Margarita Takou, Rami-Petteri Apuli, Paul DuBray, Toni Jenfors
- 2014 Teaching assistant  
Basics in genetics, 15 h

## SELECTED PRESENTATIONS

- 11/2020 Invited talk, University of Oulu, Finland  
What can the genetic and genomic architecture of adaptation tell about evolution?
- 1/2020 Oral presentation, PAG, San Diego, California  
Gene expression modularity reveals footprints of polygenic adaptation in *Theobroma cacao*
- 8/2019 Poster presentation, ESEB, Turku, Finland  
Gene expression modularity reveals footprints of polygenic adaptation in *Theobroma cacao*
- 8/2018 Oral presentation, Midwest PopGen meeting, Saint Paul, Minnesota  
Local adaptation under gene flow: Recombination, conditional neutrality and genetic trade-off shape genomic patterns in *Arabidopsis lyrata*
- 7/2018 Poster presentation, SMBE, Yokohama, Japan  
Local adaptation under gene flow: Recombination, conditional neutrality and genetic trade-off shape genomic patterns in *Arabidopsis lyrata*
- 2/2018 Oral presentation, Arabis symposium, Cologne, Germany  
Genomic patterns of local adaptation under gene flow in *Arabidopsis lyrata*
- 7/2017 Oral presentation, SMBE, Austin, Texas  
Transcriptomics and epigenetics in locally adapted *Arabidopsis lyrata*
- 8/2015 Poster presentation, ESEB, Lausanne, Switzerland  
Effects of genetic divergence on hybrid germination between subspecies of *Arabidopsis lyrata*

## REFERENCES

- Peter Tiffin (postdoctoral supervisor)  
Professor at University of Minnesota Twin Cities, USA, ptiffin@umn.edu
- Outi Savolainen (PhD supervisor)  
Emerita professor at University of Oulu, Finland, outi.savolainen@oulu.fi

## OTHER

Journal referee ([Publons](#)):

eLife, Evolution, Evolutionary Applications, Evolutionary Ecology, Frontiers in Plant Science, G3: Genes | Genomes | Genetics, Molecular Biology and Evolution, Molecular Ecology, New Phytologist, PLOS Genetics, PNAS

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