CURRICULUM VITAE
6/2021

PERSONAL DETAILS

Name	Tuomas Hämälä
Nationality	Finnish
Language	Finnish (native), English (fluent)
Email address	tuomas.hamala@gmail.com
Public profiles	Google Scholar, ResearchGate
ORCIDiD	0000-0001-8306-3397
Webpage	thamala.github.io

EMPLOYMENT

8/2018 – present — University of Minnesota Twin Cities, USA
Postdoctoral associate
Supervisor: Peter Tiffin

EDUCATION

1/2014 – 7/2018 — University of Oulu, Finland
PhD in evolutionary and population genetics
Thesis title: "Ecological genomics in *Arabidopsis lyrata*: local adaptation, phenotypic differentiation, and reproductive isolation"
Supervisor: Outi Savolainen

9/2011 – 12/2013 — University of Oulu, Finland
MSc in genetics

9/2008 – 5/2011 — University of Oulu, Finland
BSc 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**[*,†], 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€ (accepted in part)

2019   MPGI, 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   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](#)