# CURRICULUM VITAE 8/2022

# PERSONAL DETAILS

Name Tuomas Hämälä

Nationality Finland

Email address tuomas.hamala@gmail.com

Public profiles <u>Google Scholar</u>, <u>ResearchGate</u>

ORCID iD <u>0000-0001-8306-3397</u>

Webpage <u>thamala.github.io</u>

**EMPLOYMENT** 

4/2022 – present University of Nottingham, UK

Marie Skłodowska-Curie individual fellow

Supervisor: Levi Yant

9/2021 – 3/2022 University of Oulu, Finland

Academy of Finland postdoctoral fellow

8/2018 – 7/2021 University of Minnesota Twin Cities, USA

Postdoctoral associate Supervisor: Peter Tiffin

**EDUCATION** 

1/2014 – 7/2018 University of Oulu, Finland

PhD in population and evolutionary 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**

- \*equal contribution, #corresponding author
- 11. Epstein B, Burghardt L, Heath K, Grillo M, Kostanecki A, **Hämälä T**, Young N & Tiffin P<sup>#</sup> (2022). Combining GWAS and population genomic analyses to characterize coevolution in a legume-rhizobia symbiosis. *Molecular Ecology* in press.
- 10. **Hämälä T**<sup>#</sup>, 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*. *bioRxiv* preprint: 485516.
- 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<sup>#</sup> (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**<sup>#</sup> & 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**<sup>#</sup>, Guiltinan MG, Marden JH, Maximova S, dePamphilis C & Tiffin P<sup>#</sup> (2020). Gene expression modularity reveals footprints of polygenic adaptation in *Theobroma cacao*. *Molecular Biology and Evolution* 1: 110–123.
- 4. **Hämälä T**<sup>#</sup> & Savolainen O<sup>#</sup> (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.
- 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 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 PGDP, 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, transcriptome, 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

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

8/2022	Oral presentation, ESEB, Prague, Czech Republic
1/2022	Oral presentation, PopGroup55, Online
11/2021	Invited online talk, Agricultural Genomics Institute at Shenzhen, China
11/2020	Invited online talk, University of Oulu, Finland
1/2020	Oral presentation, PAG, San Diego, California
8/2019	Poster presentation, ESEB, Turku, Finland
8/2018	Oral presentation, Midwest PopGen meeting, Saint Paul, Minnesota
7/2018	Poster presentation, SMBE, Yokohama, Japan
2/2018	Oral presentation, Arabis symposium, Cologne, Germany
7/2017	Oral presentation, SMBE, Austin, Texas
8/2015	Poster presentation, ESEB, Lausanne, Switzerland

# REFERENCES

Levi Yant (postdoctoral supervisor)

Professor at University of Nottingham, UK, levi.yant@nottingham.ac.uk

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, Molecular Ecology Resources, New Phytologist, PLOS Genetics, PNAS

Programs and scripts written for NGS data analysis: GitHub