

Dr. Milan Malinsky

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EMPLOYMENT HISTORY

*SOPHiA GENETICS, Rolle, Switzerland***Algorithm Development Expert****Mar. 2025 – now**

- Algorithms to detect Copy Number Variation and Structural Variants in clinical settings (oncology, rare and hereditary disease)

*Institute of Ecology and Evolution, University of Bern, Switzerland***Principal Investigator, SNSF Ambizione Grant****Apr. 2021 – Mar. 2025**

- "Evolution of meiotic recombination and genomic divergence in percomorph fishes"

*Zoological Institute, University of Basel, Switzerland***EMBO Postdoctoral Fellowship + SNSF postdoc****Oct. 2016 – Mar. 2021**

- Advisor: Prof. Walter Salzburger
- Initially funded by a personal EMBO fellowship, this then expanded into Prof. Salzburger's SNSF grant "The genomic basis of divergence and convergence in fish species-flocks"

*Gurdon Institute, University of Cambridge, UK***Postdoctoral scientist****Oct. 2015 – Sep. 2016**

- Advisor: Prof. Eric Miska; continuing work on topics arising from my PhD

EDUCATION

*Sanger Institute, University of Cambridge, UK***PhD in Mathematical Genomics****Oct. 2011 - Sep. 2015**

- PhD defense 9th November 2015**
- Main advisor: Prof. Richard Durbin, co-advisor: Prof. Eric Miska
- Thesis: "Genomic diversity and speciation in East African cichlid fish"

*Department of Applied Mathematics, University of Cambridge, UK***Master's (MPhil) in Computational Biology****Oct. 2010 - Aug. 2011****Graduated with a distinction**

- Course covering various approaches to working with quantitative biological data, from computational neuroscience, to systems biology, to statistical genetics. Dissertation: "*C. elegans* chromatin proteins HPL-2 and LIN-13"

*University of Birmingham, UK***BSc Computer Science with Business Management, Class: I****Sep. 2007-Jun. 2010**

- Two thirds Computer Science, one third Business Management
- Final year project in the field of algorithms for medical imaging

Continuing education:**EMBO Laboratory Leadership Course, Heidelberg, Germany****Sep. 2018**

- A three-day course covering e.g. leadership roles, team development, delegation, communication, feedback + coaching, negotiation, recruitment

Workshop on Molecular Evolution, MBL, Woods Hole, MA, USA**Jul. 2013**

TEACHING ACTIVITIES

Workshops on Population and Speciation Genomics Feb. 2016, 2018, 2020; Jun. 2022

Co-organiser of two-week courses for ~70 scientists across all seniority levels

Running classes e.g., on UNIX introduction, population structure/size, probability and likelihood.

<https://evomics.org/2022-workshop-on-population-and-speciation-genomics-cesky-krumlov/>

Evolutionary Genomics MSc Course, University of Bern**May 2022 - 2024**

Development of curriculum, computer labs, lecturing on topics at the intersection of phylogenomics and population genetics

Phylogenomics Workshop - Physalia Courses**Nov. 2020**

Instructor on a one-week online course for ~25 scientist

APPROVED RESEARCH PROJECTS

"Evolution of meiotic recombination and genomic divergence in percomorph fishes",

SNSF Ambizione Grant (954,789CHF ~ €950,000), Apr. 2021 – Mar. 2025

Grant number: PZ00P3_193464; <https://p3.snf.ch/project-193464>

"Molecular basis of convergent evolution between cichlid fishes of lakes Malawi and Tanganyika",

EMBO Postdoctoral fellowship (157,659CHF ~ €155,000), Oct. 2016 – Sep. 2018

Grant number: ALTF 456-2016

"Genotype/phenotype relationships in fast evolving vertebrate species - African Lake cichlids as a model system",

Wellcome Trust PhD studentship (£155,658 ~ €185,000), Oct. 2011 – Sep. 2015

Grant number: 097677/Z/11/Z; <https://app.dimensions.ai/details/grant/grant.3640527>

SUPERVISION OF JUNIOR RESEARCHERS

Formal supervisor role:**PhD project at University of Bern – main supervisor****Sep. 2021 – now**

Marion Talbi – "Evolution of recombination landscapes"

PhD student, University of Antwerp – co-supervisor**Jan. 2021 – now**

Sophie Gresham; main PI Prof. Hannes Svandal

"The role of gene-flow in adaptive radiations"

Master's project, University of Bern**Sep. 2023 – now**

Laura Dekker – "Building a 'pangenome' for a massive adaptive radiation"

Master's project, University of Cambridge**Mar. 2013 – Jul. 2013**

Joseph J. Hanly – "Evidence for functional evolution of a miR-99b target site in related species of cichlid fishes"

Informal supervision (examples):**Training a new postdoc at the Sanger Institute****Apr. 2016**

Ian Wilson – Alignment, variant calling etc. with a high-performance computing cluster

Supervising a research assistant at the Sanger Institute**Feb. 2014 – Apr. 2014**

Daniel James – Automating genome assembly pipelines

REVIEWING ACTIVITIES

Grant proposal reviews:

European Research Council

– ERC Starting Grant, August 2022

Austrian Academy of Sciences

– DOC fellowship, May 2018

Manuscript reviews (23 manuscripts):

Current Biology

– 3x, Jan. 2017, Feb. 2019 (with W. Salzburger), May 2024

Evolution

– 1x, Dec. 2021

Genetics

– 1x, Mar. 2025

Genome Biology and Evolution

– 2x, Nov. 2015, Nov. 2021

GigaScience

– 2x, Sep. 2018 and Feb. 2019

Journal of Evolutionary Biology

– 2x, Oct. 2020, Apr. 2021

Molecular Biology and Evolution

– 2x, Jul. 2020, Oct. 2021

Molecular Ecology

– 2x, Jul. 2018, May 2024

Molecular Ecology Resources

– 2x, Aug. 2016 and Jun. 2018

Nature Ecology and Evolution

– 1x, Sep. 2018

PLoS Genetics

– 2x, Sep. 2019, Dec. 2020

Science

– 2x, Aug. 2016 (with E. A. Miska), May 2023

Systematic Biology

– 1x, Jan. 2024

Reviewing workshop applications:

Together with a panel of co-directors, selecting participants for the Workshop on Population and Speciation genomics, based on CVs and cover letters – Sep. 2017, Sep. 2019, and April 2022

PRIZES, AWARDS, FELLOWSHIPS

EMBO Postdoctoral fellowship	Oct. 2016 – Sep. 2018
Wellcome Trust PhD studentship - See "Approved research projects"	Oct. 2011 – Sep. 2015
BBSRC Master's Studentship - £18,151 (≈23,000 CHF) - Competitive studentship to fund the fees and living costs for my course in Cambridge	Oct. 2010 – Aug. 2011

RESEARCH OUTPUT

Publications:

Peer reviewed:

1. **Malinsky, M.**, Talbi, M., Zhou, Ch., Maurer, N., Sacco, S., Shapiro, B., Peichel, C. L., Seehausen, O., Salzburger, W., Weber, J. N., Bolnick, D. I., Green, R. E. & Durbin, R. (2025) Hi-reComb: constructing recombination maps from bulk gamete Hi-C sequencing, **Genetics**, iyaf150; <https://doi.org/10.1093/genetics/iyaf150>.
2. Blumer, L. M., Burskaia, V., Artiushin, I., Saha, J., Camacho Garcia, J., Elkin, J., Fischer, B., Zhou, C., Gresham, S., **Malinsky, M.**, Linderroth, T. ... (14 others) ..., Turner, G.F., Durbin, R. & Savrdal, H. (2024) Introgression dynamics of sex-linked chromosomal inversions shape the Malawi cichlid adaptive radiation. **Science** 388, eadr9961; doi: <https://doi.org/10.1126/science.adr9961>.
3. Gresham, S., Rusuwa, B., Ngochera, M., Turner, G.F., Genner, M.J., **Malinsky, M.** and Svardal, H. (2025), Genome Analyses Reveal Diverse Riverine Genetic Contributions to the Lake Malawi Cichlid Radiation. **Molecular Ecology**, e17786. <https://doi.org/10.1111/mec.17786>.
4. X García, J. I. C., **Malinsky, M.**, Joyce, D. A., Santos, M. E., Vernaz, G., Ngochera, M. J. & Svardal, H. (2025) Widespread Genetic Signals of Visual System Adaptation in Deepwater Cichlid Fishes, **Molecular Biology and Evolution**, Volume 42, Issue 7, msaf147; <https://doi.org/10.1093/molbev/msaf147>.
5. Talbi, M., Turner, G. F & **Malinsky, M.** (2024) Rapid evolution of recombination landscapes during the divergence of cichlid ecotypes in Lake Masoko. **Evolution**, qpae169, <https://doi.org/10.1093/evolut/qpae169>.
6. Sommer-Trembo, C., Santos, M. E., Clark, B., Werner, M., Fages, A., Matschiner, M., Hornung, S., Ronco, F., Oliver, C., Garcia, C., Tschopp, P., **Malinsky, M.*** & Salzburger, W.* (2024) The genetics of niche-specific behavioral tendencies in an adaptive radiation of cichlid fishes. **Science** 384, 470–475. doi: <https://doi.org/10.1126/science.adj9228>. *joint corresponding authors
7. Koppetsch, T., **Malinsky, M.** & Matschiner, M. (2024) Towards reliable detection of introgression in the presence of among-species rate variation. **Systematic Biology** syae028. doi: <https://doi.org/10.1093/sysbio/syae028>.
8. Omelchenko, D., Bitja-Nyom, A. R., Matschiner, M., **Malinsky, M.**, Indermaur, A., Salzburger, W., Bartoš, O. & Musilova, Z. (2024) Hemoglobin gene repertoire in teleost and cichlid fishes shaped by gene duplications and genome rearrangements. **Molecular Ecology**. 33:e17559. <https://doi.org/10.1111/mec.17559>.
9. Choo, L.Q., Spaggiardi, G., **Malinsky, M.**, Choquet, M., Goetze, E., Hoarau, G. & Peijnenburg, K.T.C.A. (2023), Genome-wide phylogeography reveals cryptic speciation in the circumglobal planktonic calcifier *Limacina bulimoides*. **Molecular Ecology** 32, 3200–3219. doi: <https://doi.org/10.1111/mec.16931>.
10. Barth, J.M.I., Handley, S.A., Kintzl, D., Leonard, G., **Malinsky, M.**, Matschiner, M., Meyer, B. S., Salzburger, W., Stefka, J., Trucchi, E. (2023) The history and organization of the Workshop on Population and Speciation Genomics. **Evolution: Education & Outreach** 16, 2. doi: <https://doi.org/10.1186/s12052-023-00182-w>.
11. Vernaz, G., Hudson, A. G., Santos, M. E., Fischer, B., Carruthers, M., Shechonge, A. H., Gabagambi, N. P., Tyers, A. M., Ngatunga, B. P., **Malinsky, M.**, Durbin, R., Turner, G. F., Genner, M. J. & Miska, E. A. (2022) Epigenetic Divergence during Early Stages of Speciation in an African Crater Lake Cichlid Fish. **Nature Ecology & Evolution**, 1-12. doi: <https://doi.org/10.1038/s41559-022-01894-w>.
12. Ciezarek, A., Ford, A.G.P., Etherington, G.J., Kasozic, N., **Malinsky, M.**, Mehta, T., Penso-Dolfine, L., Ngatunga, B.P., Shechonge, A., Tamatamah, R., Haerty, W., Di Palma, F., Genner, M.J. & Turner, G.F. (2022) Whole genome resequencing data enables a targeted SNP panel for conservation and aquaculture of *Oreochromis* cichlid fishes. **Aquaculture**, 548 (2). doi: <https://doi.org/10.1016/j.aquaculture.2021.737637>.
13. Vernaz, G., **Malinsky, M.**, Svardal, M., Du, M., Tyers, A. M., Santos, M. E., Durbin, R., Genner, M. J., Turner, G. F. & Miska, E. A. (2021) Mapping Epigenetic Divergence in the Massive Radiation of Lake Malawi Cichlid Fishes. **Nature Communications**, 12, 5870. doi: <https://doi.org/10.1038/s41467-021-26166-2>.

14. Svoldal, H., Salzburger, W. & **Malinsky, M.** (2021) Genetic Variation and Hybridization in Evolutionary Radiations of Cichlid Fishes. *Annu. Rev. Anim. Biosci.* 9, 55–79. doi: <https://doi.org/10.1146/annurev-animal-061220-023129>.
15. **Malinsky, M.**, Matschiner, M. & Svoldal, H. (2021) Dsuite - Fast D-statistics and related admixture evidence from VCF files. *Molecular Ecology Resources*, 21, 584–595. doi: <https://doi.org/10.1111/1755-0998.13265>.
16. Ronco, F., Matschiner, M., Böhne, A., Boila, A., Büscher, H.H., El Taher, A., Indermaur, A., **Malinsky, M.**, Ricci, V., Kahmen, A., Jentoft, S. & Salzburger, W. (2021) Drivers and dynamics of a massive adaptive radiation in cichlid fishes. *Nature*, 589, 76–81. doi: <https://doi.org/10.1038/s41586-020-2930-4>.
17. Rhie, A., McCarthy, S. A. ... (44 others) ... **Malinsky, M.**, ... (72 others) ... Durbin, R., Phillippy, A. M. & Jarvis, E. D. (2021) Towards complete and error-free genome assemblies of all vertebrate species. *Nature*, 592, 737–746. doi: <https://doi.org/10.1038/s41586-021-03451-0>.
18. Patil, C., Sylvester, J. B., Abdilleh, K., Norsworthy, M. W., Pottin, K., **Malinsky, M.**, Bloomquist, R. F., Johnson, Z. V., McGrath, P. T. & Streelman, J. T. (2021) Genome-enabled inference of evolutionary divergence in brains and behaviour. *Scientific Reports*, 11, 13016. doi: <https://doi.org/10.1101/2020.10.12.336453>.
19. Carleton, K. L., Conte, M. A., **Malinsky, M.**, Nandamuri, S. P., Sandkam, B. A., Meier, J. I., Mwaiko, S., Seehausen, O. & Kocher, T. D. (2020) Movement of transposable elements contributes to cichlid diversity. *Molecular Ecology*, 29, 4956–4969. doi: <https://doi.org/10.1111/mec.15685>.
20. Svoldal, H., Quah, F. S., **Malinsky, M.**, Ngatunga, B. P., Miska, E. A., Salzburger, W., Genner, M. J., Turner, G. F., Durbin, R. (2020) Ancestral hybridisation facilitated species diversification in the Lake Malawi cichlid fish adaptive radiation. *Mol. Biol. Evol.* 37, 1100–1113. doi: <https://doi.org/10.1093/molbev/msz294>.
21. **Malinsky, M.**, Svoldal, H., Tyers, A. M., Genner, M. J., Turner, G. F. Miska, E. A., Durbin, R. (2018) Whole genome sequences of Malawi cichlids reveal multiple radiations interconnected by gene flow. *Nature Ecology & Evolution* 2, 1940–1955. doi: <https://doi.org/10.1038/s41559-018-0717-x>.
22. **Malinsky, M.**, Trucchi, E., Lawson, D. J. & Falush, D. (2018) RADpainter and fineRADstructure: Population Inference from RADseq data. *Mol. Biol. Evol.* 35, 1284–1290. doi: <https://doi.org/10.1093/molbev/msy023>.
23. **Malinsky, M.** and Salzburger, W. (2016) Environmental context for understanding the iconic adaptive radiation of cichlid fishes in Lake Malawi. *PNAS*, 113, 11654–11656. doi: <https://doi.org/10.1073/pnas.1614272113>.
24. **Malinsky, M.**, Challis, R., Tyers, A. M., Schiffels, S., Terai, Y., Ngatunga, B. P., Miska, E. A., Durbin, R., Genner, M. J., Turner, G. F. (2015). Genomic islands of speciation separate cichlid ecomorphs in an East African crater lake. *Science* (New York, N.Y.), 350(6267), 1493-8. doi: <https://doi.org/10.1126/science.aac9927>.
25. Brawand, D., Wagner, C. E., Li, Y. I., **Malinsky, M.**, Keller, I., Fan, S., Simakov, O., Ng, A. Y., Lim, Z. W., Bezault, E., ... (76 others) ... and Di Palma, F. (2014). The genomic substrate for adaptive radiation in African cichlid fish. *Nature*, 513(7518), 375–381. doi: <https://doi.org/10.1038/nature13726>.
26. **M. Malinsky**, M. Jermyn, B. Pogue, and H. Dehghani (2010). “An Online Modeling and Image Reconstruction Tool for Optical Imaging Based on NIRFAST” in *Biomedical Optics and 3-D Imaging, OSA Technical Digest* , paper BSuD27; doi: <https://doi.org/10.1364/BIOMED.2010.BSuD27>.

Preprints:

1. Nichols, A. L. A., Shafer, M. E. R, Indermaur, A., Rüegg, A., Gonzalez-Dominguez, R., **Malinsky, M.**, Sommer-Trembo, C., Fritschi, L., Salzburger, W. & Schier, A. F. (2024) Widespread temporal niche partitioning in an adaptive radiation of cichlid fishes. *bioRxiv* 2024.05.29.596472, doi: <https://doi.org/10.1101/2024.05.29.596472>.
2. **Malinsky, M.**, Simpson, J. T. & Durbin, R. (2016) trio-sga: facilitating de novo assembly of highly heterozygous genomes with parent-child trios. *bioRxiv* 20, 051516. doi: <https://doi.org/10.1101/051516>.

Contributions to international conferences:

Conference organizer:

1. **RecomBern 2024:** “Evolution of recombination landscapes - Facilitating adaptation?”, Bern, Switzerland. www.recombern2024.org

Symposium organizer:

1. **ESEB conference** (2022) “Characterizing genomic landscapes of recombination and their evolution” Co-organized with Pierre-Alexandre Gagnaire
Link: https://www.eseb2022.cz/en/event-programme?programme_block_id=23

Oral contributions:

1. **Malinsky, M.** "Exploratory behavior and genetics of species diversification". **NWO Life (2024)**, Egmond aan Zee, Netherlands. Invited Keynote lecture.
2. **Malinsky, M.** "Genome evolution, natural selection, and recombination landscapes". **NORBIS (Norwegian Bioinformatics Graduate school) annual conference (2022)**, Rosendal (Bergen), Norway. Invited Keynote lecture.
3. **Malinsky, M.** "Evolution of genomes and meiotic recombination rates". **Biology 22 (2022)**, Basel, Switzerland.
4. **Malinsky, M.** "Evolution of recombination rate landscapes along the genome". **MACSPRO (Modelling and Analysis of Complex Systems and Processes) conference (2021)**, Moscow, Russia. Invited Keynote lecture.
5. **Malinsky, M.** "Genetic Variation in Evolutionary Radiations of Cichlid Fishes". **Cichlid science conference (2021)**, Cambridge, UK.
6. **Malinsky, M.** "Molecular basis of convergent evolution between cichlid fishes of lakes Malawi and Tanganyika". **Cichlid science conference (2019)**, Madrid, Spain.
7. **Malinsky, M., Svandal, H., Vernaz, G., Miska, E. A., Salzburger, W., Durbin, R.** "Genetic basis of functional differences between closely related species". **EMBO at Basel Life: Genomes in Biology and Medicine (2017)**, Basel, Switzerland.
8. **Malinsky, M., Svandal, H., Genner, M. J., Turner, G. F., Durbin, R., and Salzburger, W.** "Shared variation between Great Lakes: do they all use the same variation?" **Cichlid science conference (2017)**, Prague, Czech Republic.
9. **Malinsky, M., Trucchi, E., Lawson, D. J. & Falush, D.** "RADpainter and fineRADstructure: Population Inference from RADseq Data" **Society for Molecular Biology and Evolution (SMBE) annual meeting (2016)**, Brisbane, Australia.
10. **Malinsky, M., Tyers, A., Miska, E. A., Genner, M. J., Durbin, R., Turner, G. F.** "Speciation in Lake Malawi and in Tanzanian crater lakes". **Cichlid science conference (2015)**, Graz, Austria.
11. **Malinsky, M., Challis, R. J., Tyers, A., Miska, E. A., Genner, M. J., Durbin, R., Turner, G. F.** "Speciation in cichlid fishes in Lake Malawi catchment" **Society for Molecular Biology and Evolution (SMBE) annual meeting (2015)**, Vienna, Austria.
12. **Malinsky, M.** "Genomic diversity and speciation in East African cichlid fish". **Cambridge Evolutionary Genetics & Genomics Symposium (2015)**, Cambridge, UK.
13. **Malinsky, M., Challis, R. J., Tyers, A., Miska, E. A., Genner, M. J., Durbin, R., Turner, G. F.** "Islands of positive selection and speciation with gene flow in African crater lake cichlids". **Society for Molecular Biology and Evolution (SMBE) annual meeting (2014)**, San Juan, USA.

Poster presentations:

1. **Malinsky, M.** "Disentangling the genomic interplay of selection and recombination" **Society for Molecular Biology and Evolution (SMBE) annual meeting (2023)**, Ferrara, Italy.
2. **Malinsky, M.** "Disentangling the genomic interplay of selection and recombination" **European Society for Evolutionary Biology (ESEB) congress (2022)**, Prague, Czech Republic.
3. **Malinsky, M. and Matschiner, M.** "c-genie - Assessing the impact of recombination on phylogenomic inference" **Society for Molecular Biology and Evolution (SMBE) annual meeting (2018)**, Yokohama, Japan.
4. **Malinsky, M. and Matschiner, M.** "microRNA evolution in East African cichlids" **Society for Molecular Biology and Evolution (SMBE) annual meeting (2013)**, Chicago, USA.
5. **Malinsky, M. and Dehghani, H.** "An Online Modeling and Image Reconstruction Tool for Optical Imaging Based on NIRFAST". **The Optical Society of America annual meeting (2010)**, Miami, USA.

General contributions to science:

1. Cambridge University Museum of Zoology - Lake Malawi cichlid fish collection
 - Key role in establishing one of the largest Lake Malawi fish collections in the world (>1200 specimens, >200 species)
 - Planning and participating in the three-week field expedition in 2016, participating in the cataloguing in the museum
 - The full collection is accessible to researchers and used in teaching in Cambridge too. See e.g.: https://youtu.be/XYyu6ln_khQ; <https://www.zoo.cam.ac.uk/news/teaching-and-the-museum-of-zoology>
2. Software package trio-sga: <https://github.com/millanek/trio-sga> (see the Publications - Preprints)
 - An extension of the SGA *de novo* genome assembly tool by Jared Simpson
 - Facilitating assembly of highly heterozygous genomes by using data from mother-father-offspring trios

3. Software package fineRADstructure: <https://github.com/millanek/fineRADstructure>
 - Widely used population structure inference package, using clustering code from fineSTRUCTURE
 - Publication in Molecular Biology and Evolution (see “Publications in peer-reviewed journals”)
4. Software package D-suite: <https://github.com/millanek/Dsuite>
 - Widely used package for detecting signatures of gene-flow among species or among populations from genomic polymorphism data
 - Publication in Molecular Ecology Resources (see “Publications in peer-reviewed journals”)
5. Co-author, with Michael Matschiner of the c-genie software: <https://github.com/millanek/c-genie>
 - Simulation-based software for assessing the impact of recombination on phylogenomic inference
 - Presented by me at the SMBE annual meeting in 2018
6. Generated two high quality *de novo* genome assemblies from long read (PacBio) data; now publicly available:
 - *Astatotilapia calliptera* - http://www.ensembl.org/Astatotilapia_calliptera/Info/Index
 - *Simochromis diagramma* - https://www.ncbi.nlm.nih.gov/assembly/GCA_900408965.1/
7. microRNA gene annotation for five cichlid fish genomes
 - e.g., <http://www.mirbase.org/summary.shtml?org=mze>

REFEREES

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