Kimberly J. Gilbert | CV

Computational and Molecular Population Genetics Lab
Institute of Ecology and Evolution – University of Bern
Baltzerstrasse 6, 3012 Bern, Switzerland

kimberly.gilbert@iee.unibe.ch • http://kjgilbert.github.io/

Population geneticist & evolutionary biologist

Employment

EMBO Post-doctoral Fellow — **2017 – Present**
University of Bern — *Bern, Switzerland*
Advisor: Prof. Dr. Laurent Excoffier

EEB Post-doctoral Fellow — **2016 – 2017**
University of Toronto — *Toronto, Ontario*
Advisors: Dr. Aneil F. Agrawal, Dr. Stephen I. Wright

Education

Ph.D. Zoology — **Sep 2011 – Oct 2016**
University of British Columbia — *Vancouver, British Columbia*
Advisor: Dr. Michael C. Whitlock
Dissertation title: Understanding local adaptation and effective population size in the face of complex demographic history

B.Sc. Biology — **2006 – 2010**
University of Virginia — *Charlottesville, Virginia*
Graduated with Distinction
Specialization in environmental & biological conservation

Publications

13. Peischl S, Dupanloup I, Foucal A, Jomphe M, Bruat V, Grenier J-C, Gouy A, **Gilbert KJ**, Gbeha E, Bosshard L, Hip-Ki E, Agbessi M, Hodgkinson A, Vézina H, Awadalla P, Excoffier L (2017) Relaxed selection during a recent human expansion. *Genetics*, genetics.300551.2017
12. **Gilbert KJ**, MC Whitlock. (2017) The genetics of adaptation to discrete heterogeneous environments: Frequent mutation or large effect alleles can allow range expansion. *Journal of Evolutionary Biology*, 30(3), 591-602. doi:10.1111/jeb.13029.
11. **Gilbert KJ**, NP Sharp, AL Angert, GL Conte, JA Draghi, F Guillaume, AL Hargreaves, R Matthey-Doret, MC Whitlock. (2017) Local maladaptation reduces expansion load during range expansion. *The American Naturalist*, 189(4), 368-380, doi:10.1086/690673.
10. **Gilbert KJ** (2016) Identifying the number of population clusters with STRUCTURE: Problems and solutions. *Molecular Ecology Resources*, 16(3), 601-603.
9. **Gilbert KJ**, MC Whitlock (2015) Evaluating methods for estimating local effective population size with and without migration. *Evolution*, 68(8), 2154-2166.

8. Santiso X, L Lopez, **KJ Gilbert**, R Barreiro, MC Whitlock, R Retuerto (2015) Patterns of genetic variation within and among populations in *Arbutus unedo* and its relation with selection and evolvability. *Perspectives in Plant Ecology, Evolution and Systematics*, 17(3), 185-192.
7. **Gilbert KJ**, MC Whitlock (2015) Q_{ST}-F_{ST} comparisons with unbalanced half-sib designs. *Molecular Ecology Resources*, 15(2), 262-267.
6. Caplins SA, **KJ Gilbert**, C Ciotir, J Roland, SF Matter, N Keyghobadi (2014) Landscape structure and the genetic effects of a population collapse. *Proceedings of the Royal Society B*. 281: 20141798; doi: 10.1098/rspb.2014.1798
5. Vines TH, AYK Albert, RL Andrew, F Débarre, DG Bock, MT Franklin, **KJ Gilbert**, J-S Moore, S Renaut, DJ Rennison (2014) The availability of research data declines rapidly with age. *Current Biology*, 24, 94-97.
4. Vines TH, RL Andrew, DG Bock, MT Franklin, **KJ Gilbert**, NC Kane, EJ Kleynhans, J-S Moore, BT Moyers, S Renaut, DJ Rennison, T Veen, S Yeaman (2013) Mandated archiving greatly improves access to research data. *FASEB Journal*, 27(4), 1304-1308.
3. **Gilbert KJ**, RL Andrew, DG Bock, MT Franklin, NC Kane, J-S Moore, BT Moyers, S Renaut, DJ Rennison, T Veen, TH Vines (2012) Recommendations for utilizing and reporting population genetic analyses: The reproducibility of genetic clustering using the program STRUCTURE. *Molecular Ecology*, 21(20), 4925-4930.
2. Keller SR, **KJ Gilbert**, PD Fields, DR Taylor (2012) Bayesian inference of a complex invasion history revealed by nuclear and chloroplast genetic diversity in the colonizing plant, *Silene latifolia*. *Molecular Ecology*, 21(19), 4721-4734.
1. Whitlock MC, **KJ Gilbert** (2012) Q_{ST} in a hierarchically structured population. *Molecular Ecology Resources*, 12(3), 481–483.

Awards & Funding

EMBO Long-term Post-doctoral Fellowship **2017 – 2019**
143,400 CHF

Ecology and Evolutionary Biology Post-doctoral Fellowship **2016 – 2017**
University of Toronto *$40,500 CAD* Accepted in part

Declined **– NSF Post-doctoral Research Fellowship** **2016 – 2019**
National Plant Genome Initiative *$210,000 USD*

Cordula and Gunter Paetzold Fellowship **2015 – 2016**
University of British Columbia *$18,000 CAD*

Declined **– Zoology Graduate Fellowship** **2015 – 2016**
University of British Columbia *$16,000 CAD*

Ann and William Messenger Graduate Fellowship **2015**
University of British Columbia *$700 CAD*

Zoology Graduate Fellowship **2014 – 2015**
University of British Columbia *$11,000 CAD*

Frieda Granot Graduate Scholarship in Interdisciplinary Research **2013 – 2014**
University of British Columbia *$200 CAD*

Theodore E Arnold Fellowship **2013 – 2014**
University of British Columbia *$7,750 CAD*

Patrick David Campbell Graduate Fellowship **2013 – 2014**
University of British Columbia *$8,050 CAD*

Zoology Graduate Fellowship **2013 – 2014**
University of British Columbia *$11,000 CAD*

Zoology Graduate Student Travel Award **2013, 2014, 2016**
University of British Columbia *$500; $500; $400 CAD*

BRITE Fellowship **2011 – 2013**
University of British Columbia $21,000 CAD

Presentations & Workshops

Invited

Species range shifts and local adaptation **January 2018**
AndinA workshop *Bariloche, Argentina*
35-member international working group combining ecological and evolutionary disciplines to further understanding of range expansions and local adaptation

Local adaptation, expansion load, and mutation load **Sept., Oct. 2017**
University of Basel – Botanical Colloquium *Basel, Switzerland*
Uppsala University *Uppsala, Sweden*

The genetics of adaptation during expansion across heterogeneous environments **April 2017**
University of Zürich; Behaviour, Ecology, Environment, and Evolution Seminar Series *Zürich, Switzerland*

Local maladaptation reduces expansion load during species range expansion **July 2016**
CSEE "Theoretical ecology and evolutionary biology" symposium *St. John's, Newfoundland*

Data availability, archiving, and scientific reproducibility **June 2016**
American Society of Mammalogists Annual Conference *Minneapolis, MN*
''Big data meets mammalogy: how to find and share data" symposium

Population genetic inference in the face of demographic history **August 2015**
University of Bern *Bern, Switzerland*

Estimating effective population size and the reproducibility of science **Feb. 2015, Dec. 2014**
Monash University *Melbourne, VIC, Australia*
Duke University Pop Bio Seminar Series *Durham, NC*

Reproducible Science Hackathon **December 2014**
NESCent Working Group *Durham, NC*
21-member working group aiming to develop a curriculum and workflow for teaching reproducible science

SimBank **November 2014**
NESCent Catalysis Meeting *Durham, NC*
25-member working group aiming to create a collection of openly available simulation results to facilitate testing of statistical population genetic and phylogeographic methods

Contributed

Recovery from expansion load is limited during species range shifts **2017**
CeMEB Assembly: Biological invasions & range expansions from an evolutionary perspective *Tjärno, Sweden*

Mutation load across mating systems: how does load change and how is it best estimated **2017**
SMBE Meeting - talk *Austin, TX*
Evolution Meeting - talk *Portland, OR*

Local maladaptation reduces expansion load during species range expansion **2016**
Evolution Meeting - talk *Austin, TX*

Local adaptation and range expansions **2015**
SFU-UBC-UVic-UW Ecology and Evolution Retreat - talk *Brackendale, BC*

Validating SNP loci underlying local adaptation in lodgepole pine **2015**
15^{th} ESEB Congress - poster *Lausanne, Switzerland*

Evaluating methods to estimate effective population size in the presence of migration **2014**
Evolution Meeting - talk *Raleigh, NC*

Evo-WIBO Conference (Evolutionary Biology in the Pacific Northwest) - talk — *Port Townsend, WA*

Estimating effective population size in natural populations — **2013**
Evolution Meeting - talk — *Snowbird, UT*
SFU-UBC-UVic Ecology and Evolution Retreat - talk — *Brackendale, BC*

Effective population size estimates in a metapopulation of *Silene latifolia* — **2012**
1st Joint Congress on Evolutionary Biology - poster — *Ottawa, ON*

Range expansion and adaptation across heterogeneous environments — **2012**
Landscape Genetics Symposium, CIEE Graduate Mini-Course - talk — *Toronto, ON*
Evo-WIBO Conference (Evolutionary Biology in the Pacific Northwest) - poster — *Port Townsend, WA*

Inferred invasion history of *Silene latifolia* into North America — **2011**
13th ESEB Congress - poster — *Tuebingen, Germany*
SFU-UBC-UVic Ecology and Evolution Retreat - poster — *Brackendale, BC*

Teaching Experience

Molecular Population Genetics Practical — **Spring 2018**
University of Bern
Teaching assistant for practicals in molecular data analysis for undergraduate biology majors

Statistics for Biology — **Spring 2017**
University of Bern
Teaching assistant for three sessions of practicals in statistics for undergraduate biology majors

Fundamentals of Biostatistics — **Fall 2013, 2014, 2015**
BIOL 300 — *UBC*
Statistical procedures for biological research; estimation, hypothesis testing, goodness of fit, analysis of variance and regression; use of computers for statistical analysis
Two sections (70 students total) in 2013, one section (36 students) in 2014, one section (36 students) in 2015.
Lab coordinator for 254 students enrolled in 2014 and for 275 students enrolled in 2015.

***Guest Lecture* – Quantitative Methods in Ecology and Evolution** — **January 2013**
BIOL 548 — *UBC*
Guest lecture on making maps in R

Fundamentals of Evolutionary Biology — **Fall 2012, Spring 2013**
BIOL 336 — *UBC*
Discussion-based tutorial covers natural selection, population genetics, quantitative genetics, systematics, and classical and molecular approaches to the study of evolution
Three discussion sections of 45 students total per semester

Scientific Software

QstFstComp: R package for performing balanced and unbalanced Q_{ST}–F_{ST} analyses, available at http://github.com/kjgilbert/QstFstComp

aNEMOne: R package to serve as a wrapper, creating inputs and analyzing outputs, for the simulation program Nemo, available at http://github.com/kjgilbert/aNEMOne

Volunteer & Outreach

Subject Editor: *Molecular Ecology*, *Molecular Ecology Resources* (2018 – 2021)

Reviewer: *New Phytologist*, *Molecular Ecology*, *Molecular Ecology Resources*, *Heredity*, *Ecology and Evolution*, *PeerJ*, *Ecography*, *Tree Genetics & Genomes*

IEE Junior Staff Seminar Series **2017 – Present**
One of six junior staff organizing the invited lecture series in ecology and evolution

Symposium co-organizer **2018**
Joint 2018 ESEB-SSE-ASN-SSB meeting in Montpellier, France
Co-organized with Drs. Nathaniel Sharp, Frédéric Austerlitz, and Paul Verdu a symposium entitled "From theory to genome-wide data: inferring selection, demography, gene flow and admixture"

Graduate Student Council Member **2013 – 2016**
American Society of Naturalists
Council Chair for 2015-2016 term
Organized the student-mentor mixer at the 2016 Evolution Meeting in Austin, TX
Organized the student-mentor mixer at the 2014 Evolution Meeting in Raleigh, NC
Served on the workshops committee for ASN-sponsored workshops

Faculty Search Committee: Graduate Student Representative **2014**
Evolutionary biology CRC2 job search for the Department of Zoology, University of British Columbia

Journal club organizer
UBC Evolution Discussion Group (EDG) weekly reading group *2014 – 2015*
Evolution discussion group at University of Bern, Institute of Ecology and Evolution *2017 – Present*

Volunteer mist-netting and bird banding **2013 – 2016**
Wild Research *Iona Island Bird Observatory, Vancouver, BC*
Participate in winter, spring migration, and fall migration bird monitoring
Teach proper bird handling, aging, data collection, and mist net extraction techniques to new volunteers
Assist in teaching other volunteers and visitors to the station about the species conservation and monitoring, and the general tasks of running a banding station

Previous Research Experience

Independent Study & Research Technician **Sep. 2009 – June 2011**
University of Virginia
Genetic analysis of metapopulation processes in the *Silene-Micobotryum* host-pathogen system
Supervisors: Dr. Douglas R. Taylor, Dr. Peter D. Fields, Dr. Janis Antonovics

Field Technician & Research Assistant **May 2009 – August 2009**
Blandy Experimental Farm, University of Virginia
Field research on effects of tropospheric ozone on native vs. invasive tree species
Supervisors: Dr. David E. Carr, Dr. Eric E. Elton

MAPS Bird Banding Intern **May 2008 – August 2008**
Monitoring Avian Productivity and Survivorship – The Institute for Bird Populations
Mist-netting, banding, and processing passerines and near-passerines during summer breeding season to monitor populations of local species
Supervisor: James Junda, M.Sc.