Project Summary

Overview testing

Intellectual Merit

Broader Impacts

Project Description

Introduction

nicely written intro goes here

Objectives

Objective I: do this cool thing

talk about the objectives here, then list them specifically:

- 1. Do ...?
- 2. Do ...?
- 3. Is there evidence of ...?

Objective II: Determine the extent to which ...

Maize was domesticated in...

Rationale and Significance

rationale and significance

Research Plan

Assess the evolutionary role of ...

first subsection stuff

Does the potential for ...?

lots of subsection stuff

Is i...?

subsection stuff

Can a widespread species serve as ...?

subsection stuff

Potential Challenges potential challenges

Broader Impacts

stuff here

subsection

stuff

subsection

stuff

Results From Prior NSF Support

don't think I need this section

Biographical Sketch — Kimberly Julie Gilbert

A. Professional Preparation

	Institution, Location	Major	Degree	Year
•	University of Virginia, USA	Biology	B.Sc.	2010
	University of British Columbia, Canada	Zoology	Ph.D.	2016 (expected)
	Advisor: Michael C. Whitlock			

B. Publications

Five Publications Most Closely Related to the Proposed Project

- 1. **Gilbert KJ**, MC Whitlock (2015) Evaluating methods for estimating local effective population size with and without migration. *Evolution*, 68(8), 2154-2166.
- 2. **Gilbert KJ**, MC Whitlock (2015) *Q*st-*F*st comparisons with unbalanced half-sib designs. *Molecular Ecology Resources*, 15(2), 262-267.
- 3. 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
- 4. 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.
- 5. Whitlock MC, **KJ Gilbert** (2012) *Q*s⊤ in a hierarchically structured population. *Molecular Ecology Resources*, 12(3), 481–483.

Four Other Significant Publications

- 1. 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.
- 2. 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.
- 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.

C. Select Conference Presentations (chosen from 11 presentations)

- 2015 Validating SNP loci underlying local adaptation in lodgepole pine; KJ Gilbert, S Yeaman, KE Lotterhos, KA Hodgins, H Suren, JA Holliday, S Nadeau, SN Aitken, MC Whitlock Poster, 15th ESEB Congress, Lausanne, Switzerland
- 2014 Evaluating methods for estimating effective population size in the presence of migration; KJ Gilbert & MC Whitlock *Oral presentation, Evolution, Raleigh, USA*
- 2012 Range expansion and adaptation across heterogeneous environments; KJ Gilbert & MC Whitlock Poster, Evo-WIBO Conference (Evolutionary Biology in the Pacific Northwest), Port Townsend, USA
- 2011 Inferred invasion history of *Silene latifolia* into North America utilizing population genetic data and approximate Bayesian computation; KJ Gilbert, SR Keller, PD Fields, DR Taylor *Poster,* 13th Congress of the European Society for Evolutionary Biology, Tuebingen, Germany

D. Grants and Awards

Cordula and Gunter Paetzold Fellowship, UBC \$18,000CAD

2015 - 2016





Declined; Zoology Graduate Fellowship, UBC \$11,000CAD	2015 - 2016
Ann and William Messenger Graduate Fellowship, UBC \$700CAD	2015
Zoology Graduate Fellowship, UBC \$11,000CAD	2014 – 2015
Frieda Granot Graduate Scholarship in Interdisciplinary Research \$200CAD	2013 - 2014
Theodore E Arnold Fellowship \$7,750CAD	2013 - 2014
Patrick David Campbell Graduate Fellowship \$8,050CAD	2013 - 2014
Declined; Zoology Graduate Fellowship, UBC \$10,000CAD	2013 - 2014
BRITE Fellowship, UBC \$10,500CAD per annum	2011 – 2013

E. Synergistic Activities

- 1. Working Groups: Participated in the NESCent Reproducible Science Hackathon (2014), a 21-member working group aimed at developing a curriculum and workflow for teaching reproducible science to researchers of any background. Participated in the SimBank NESCent Catalysis Meeting (2014) which was a 25-member working group to create a collection of openly available simulation results to facilitate testing of statistical population genetic and phylogeographic methods.
- 2. Teaching: Teaching assistant for Fundamentals of Evolutionary Biology (Fall 2012-Spring 2013) where I taught three sections per term of 45 students each and lead discussion-based tutorials. Teaching assistant for Fundamentals of Biostatistics (Fall 2013, 2014, 2015). Taught two sections of 70 students total in 2013, and in 2014 and 2015 served as the lab coordinator for 254 and 276 students enrolled in the course, respectively, while teaching one section of 36 and 35 students respectively.
- 3. Service: Served as a Graduate Student Council Member for the American Society of Naturalists (2013-2016, chair 2015-2016). Served as the graduate student representative on the 2014 evolutionary biology CRC2 job search for the Department of Zoology, University of British Columbia. Organize the Biodiversity Research Centre's weekly evolution discussion group (2014-2016), for students, post-docs, and faculty from the departments of Zoology, Botany, Forestry, and Fisheries to discuss current papers in evolutionary biology. Reviewer for Molecular Ecology Resources, Ecology and Evolution, Tree Genetics & Genomes.
- 4. Outreach: Volunteer mist-netting and bird banding with local Vancouver non-profit organization Wild Research (2013-2015) where I participated in winter, spring migration, and fall migration bird monitoring at Iona Island Bird Observatory, taught volunteers proper bird handling, aging, data collection, and mist net extraction techniques, and assisted in educating public visitors to the station about the species conservation and monitoring, and the general tasks of running a banding station.

Biographical Sketch — Jeffrey Ross-Ibarra

1 Professional Preparation

Institution	Area	Degree / Training	Dates
University of California Riverside	Botany	BA, MS	1998, 2000
University of Georgia	Genetics	PhD	2006
University of California Irvine	Genetics	Postdoctoral Research	2008

2 Professional Appointments

Position	Institution	Dates
Associate Professor	University of California Davis	2012-present
Assistant Professor	University of California Davis	2009-2012
Profesor de Asignatura	Universidad Nacional Autónoma de México	2001

3 Products

Most Relevant to the Proposed Research

- Mezmouk S, Ross-Ibarra J (2014) The pattern and distribution of deleterious mutations in maize. (2014) G3 4:163-171
- Hufford MB, Xun X, van Heerwaarden J, Pyhäjärvi T, Chia J-M, Cartwright RA, Elshire RJ, Glaubitz JC, Guill KE, Kaeppler S, Lai J, Morrell PL, Shannon LM, Song C, Spinger NM, Swanson-Wagner RA, Tiffin P, Wang J, Zhang G, Doebley J, McMullen MD, Ware D, Buckler ES, Yang S, Ross-Ibarra J (2012) Comparative population genomics of maize domestication and improvement. NATURE GENETICS 44:808-811
- Cook JP, McMullen MD, Holland JB, Tian F, Bradbury P, Ross-Ibarra J, Buckler ES, Flint-Garcia SA (2012) Genetic architecture of maize kernel composition in the Nested Association Mapping and Inbred Association panels. PLANT PHYSIOLOGY 158: 824-834
- van Heerwaarden J, Doebley J, Briggs WH, Glaubitz JC, Goodman MM, Sánchez González JJ, **Ross-Ibarra J** (2011) Genetic signals of origin, spread and introgression in a large sample of maize landraces. PNAS 108: 1088-1092
- Ross-Ibarra J, Tenaillon M, Gaut BS (2009) Historical divergence and gene flow in the genus Zea. GENETICS 181: 1399-1413.

Additional Products

- Gerke JP, Edwards JW, Guill KE, Ross-Ibarra J, McMullen MD (2015) The genomic impacts
 of drift and selection for hybrid performance in maize. GENETICS In Press
- Takuno S, Ralph P, Swarts K, Elshire RJ, Glaubitz JC, Buckler ES, Hufford MB, and Ross-Ibarra J (2015) Independent molecular basis of convergent highland adaptation in maize. GENETICS 200:1297-1312
- Wills DM, Whipple C, Takuno S, Kursel LE, Shannon LM, **Ross-Ibarra J**, Doebley JF (2013) From many, one: genetic control of prolificacy during maize domestication. PLoS GENETICS 9(6): e1003604.
- Studer A, Zhao Q, **Ross-Ibarra J**, Doebley J (2011) Identification of a functional transposon insertion in the maize domestication gene *tb1*. NATURE GENETICS 43:1160-1163.

 Gore MA, Chia JM, Elshire RJ, Sun Q, Ersoz ES, Hurwitz BL, Peiffer JA, McMullen MD, Grills GS, Ross-Ibarra J, Ware DH, Buckler ES (2009) A first-generation haplotype map of maize. SCIENCE 326: 1115-1117.

4 Synergistic Activities

- Faculty Development Award in recognition of university service, 2015
- Editor, G3, PeerJ, Axios Reviews
- DuPont Young Professor 2012-2014 and faculty advisor DuPont Pioneer graduate student symposium in plant breeding 2012-present
- Functional Genetics of Maize Centromeres US-Mexico exchange program, 2011-present
- Presidential Early Career Award for Scientists and Engineers 2009

5 Collaborators and Other Affiliations

Collaborators and Co-editors (Total: 56)

Cornell U Peter Bradbury, Jeffrey Glaubitz, Susan McCouch, Qi Sun, Feng Tian, Sharon Mitchell; USDA-ARS Edward Buckler, Sarah Hake, James Holland, Sherry Flint-Garcia, Mike McMullen, Doreen Ware, Jode Edwards; U Southern California Peter Ralph; UC Davis Alan Bennet, Daniel Runcie, Ed Taylor, Graham Coop, Keith Bradnam, Ian Korf, David Neale, Amélie Gaudin; UC Irvine Kevin Thornton; Carnegie Institute Davide Sosso; Stanford Wolf Frommer; LANGEBIO Ruairidh Sawers; U Georgia Kelly Dawe; Arizona State Reed Cartwright; U Missourri James Birchler, Katherine Guill, David Wills; Beijing Genomics Institute Song Chi, Xun Xu; U Wisconsin John Doebley, Jiming Jiang, Shawn Kaeppler; Syngenta William Briggs; Monsanto Lisa Kanizay; Dupont Pioneer Andy Baumgarten, Justin Gerke, Oscar Smith, Tabare Abadie; U Minnesota Roman Briskine, Peter Morrell, Chad Myers, Nathan Springer, Peter Tiffin; MIT Mary Gehring; NC State Major Goodman; INRA Clementine Vitte, Maud Tenaillon; Brigham Young Clinton Whipple; Danforth Center Anthony Studer; Universidad de Guadalajara Jesus Sànchez Gonzàlez; Iowa State Carolyn Lawrence; U Hawaii Gernot Presting; UC Riverside Mitchell Provance

Graduate Advisors and Postdoctoral Sponsors (Total: 3)

UC Riverside Norman Ellstrand; U Georgia James Hamrick; UC Irvine Brandon Gaut

Thesis Advisor and Postgraduate Sponsor (Total: 14)

Postdoctoral: *Iowa State* Matthew Hufford; *Graduate U Advanced Studies* Shohei Takuno; *U Oulu* Tanja Pyhäjärvi, *KWS* Sofiane Mezmouk; *Wageningen* Joost van Heerwaarden; *USDA* Tim Beissinger; *UC Davis* Kate Crosby, Sayuri Tsukahara, Simon Renny-Byfield, Jinliang Yang **Graduate:** Dianne Velasco, Paul Bilinski, Anna O'Brien, Michelle Stitzer

Biographical Sketch — Kevin Richard Thornton

6 Professional Preparation

Institution	Area	Degree / Training	Dates
University of Puget Sound	Botany	BA	1997, 2000
University of Chicago	Genetics	PhD	2003
Cornell University	Genetics	Postdoctoral Research	2007

7 Professional Appointments

Position	Institution	Dates
Associate Professor	University of California Irvine	2012-present
Assistant Professor	University of California Irvine	2007-2012

8 Products

Most Relevant to the Proposed Research

- **Thornton**, K. R. (2014) A C++ template library for efficient forward-time population genetic simulation of large populations. Genetics 98:157-166 PMID: 24950894
- Thornton, K. (2003) libsequence, a C++ class library for evolutionary genetic analysis. Bioinformatics 19(17): 2325-2327 PMID 14630667
- Thornton, K. R., A. J. Foran, and A. D. Long (2013) Properties and modeling of GWAS when complex disease risk is due to non-complementing, deleterious mutations in genes of large effect. PLoS Genetics 9: e1003258. PMID 23437004

Additional Products

- Cridland, J. M., K. R. Thornton and A. D. Long (2015) Gene expression variation in Drosophila melanogaster due to rare transposable element insertion alleles of large effect. Genetics 199: 85-93.
- Baldwin-Brown, J., A. D. Long, and K. R. Thornton (2014) The Power to Detect Quantitative Trait Loci Using Resequenced, Experimentally Evolved Populations of Diploid, Sexual Organisms. Molecular Biology and Evolution 31: 1040-1055. PMID 24441104
- Open-source software: http://molpopgen.github.io/fwdpp/
- Open-source software: http://molpopgen.github.io/libsequence/

9 Synergistic Activities

- Open-source software: http://molpopgen.github.io/fwdpy/ This software is unpublished, and will be a key resource for this proposal.
- Editor, G3

10 Collaborators and Other Affiliations

Collaborators and Co-editors (Total: 7)

- Cornell University Andrew G. Clark
- North Carolina State University Trudy Mackay
- Princeton University Peter Andolfatto
- Rochester University Daniel Garrigan, Daven C. Presgraves
- UC Irvine Anthony (Tony) Long
- University of Kansas Stuart MacDonald

Graduate Advisors and Postdoctoral Sponsors (Total: 2)

University of Chicago Manyuan Long Cornell University Andrew G. Clark

Thesis Advisor and Postgraduate Sponsor (Total: 3)

Postdoctoral: Rebekah R. Rogers PhD Thesis Advisor: Julie M. Cridland, Jaleal S. Sanjak

Data Management Plan

Data Types

This proposal will generate genotype and full-genome sequence data, phenotype data, analytical code, germplasm, and publications.

Data Archiving, Plan for Sharing, Public Access Policy

Genotype and Sequence Data EDIT THIS WHOLE DOCUMENT

Dissertation Summary - Kimberly J. Gilbert

A major obstacle in evolutionary biology is the difficulty of population genetic inference in the face of confounding factors, such as demographic history. My dissertation work has focused on several topics related to this broader area of research:

- 1. Estimating effective population sizes from genetic data in the face of migration
- 2. Evaluating the factors related to species' abilities to locally adapt at range edges during expansion
- 3. Validating SNP loci under selection in lodgepole pine (*Pinus contorta*) for local adaptation to climate

talk about NE work talk about range edge stuff talk about adaptree stuff