

METE K. YUKSEL

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EDUCATION

Ph.D. in Ecology and Evolutionary Biology (EEB)

September 2022 - present | University of Toronto (UT), Toronto, Ontario

Supervisors: Drs. Matt Osmond and Nicole Mideo

B.S. in Mathematics *General and Applied Quantitative Modeling Options*

Major GPA: 4.00

B.S. in Statistical Science *General Option*

Major GPA: 4.00

August 2018 - May 2022 | University of Idaho (UI), Moscow, ID

Cumulative GPA: 3.92

Relevant Coursework:

- Proof via number theory
- Linear algebra
- **Ordinary* & partial* differential equations**
- **Mathematical biology**
- Microbiology
- Cell & molecular biology
- Systems biology
- **Genomics**
- **Mathematical genetics***
- Evolutionary biology*
- **Molecular evolution***
- **Bias in STEM***
- **R programming**
- Modern physics
- Probability theory*
- Mathematical statistics
- Stochastic processes
- **Theory of linear models***
- **Bayesian statistics***
- **Computational statistics***
- Sample survey methods
- **Likelihood inference***
- Point-set topology*
- Advanced calculus
- Measure & integration theory*
- Complex analysis*
- Fourier analysis*
- Functional analysis*
- **Numerical methods**
- Groups & fields*
- Commutative algebra

* indicates graduate-level coursework. **Bolded** courses are of particular relevance. **Blue** text indicates course was taken at the University of Toronto; otherwise, courses were taken at the University of Idaho.

RESEARCH INTERESTS & EXPERIENCE

Interests: host-parasite interactions; evolutionary genetics and ecology of zoonotic emergence; recombination rate variation; coalescent theory & phylodynamics.

EVOLUTION OF THE RECOMBINATION MAP

Department of Ecology & Evolutionary Biology, University of Toronto | September 2023 --- present

- Developing mathematical framework to understand how the distribution of recombination breakpoints evolves, under the constraint of a total genome-wide rate. The project is motivated by the observation that the location of recombination breakpoints alters the effective rate of recombination: breaks near the center of the chromosome shuffle, on average, more alleles than do breaks near the tips.
- The model describes the evolution of the recombination landscape evolves as deleterious mutations and "local" modifiers accumulate along individual chromosomes.

SPILLCOMBINA VOIR: RECOMBINATION AND VIRAL EMERGENCE

Department of Ecology & Evolutionary Biology, University of Toronto | September 2023 --- present

- Developed theory to understand how recombination shapes pathogen evolution in wildlife and domestic animal reservoirs, and the spillover and sustained transmission of such pathogens in human hosts.
- Modeling shows that the extent of recombination between loci underlying emergence is greatest in short-lived, acutely infected hosts and the extent of linkage disequilibrium (i.e., statistical associations between such mutations) greatest at the other end of the life history axis.
- We test predictions of the theory using all avian H5Nx sequences from 2022-present. We do this in two ways. First, we calculate linkage disequilibrium between mutations found on different segments of the flu genome and regress mean LD on body size, a proxy for mean lifetime. The results are consistent with predictions of the theory. Second, we infer reassortment networks and rates for all segment pairs (e.g., the HA and NA, PB1 and MP) associated to a host (e.g., mallards) and flyway (e.g., East of the Rockies).
- Publication in prep.

MODELING TO INFORM MANAGEMENT OF CHINOOK SALMON IN THE WILLAMETTE

Department of Fish and Wildlife Sciences, University of Idaho | May 2022 --- August 2022

- With Matt Keefer and Dr. Chris Caudill, analyzed a statistical model of Chinook Salmon abundance (with life history-specific mortality, environmental stochasticity, etc.) in the Willamette River system.
- Built a Shiny app to inform management decisions made by the U.S. Army Corps of Engineers.
- Technical report written and shared with US Army Corps. Publication in prep.

STATISTICAL (EPI)GENETICS OF BREAST CANCER

- Inspired by a previous study using gene expression data from various breast and blood cancer sub-types, a collaborator and I compiled, cleaned, and used Illumina 450K methylation array data to
 1. identify differentially methylated genes between healthy, stem cell, and breast cancer tissues
 2. reconstruct the phylogenetic relationships between the tissue types.

COMPETITIVE EXCLUSION BY CURVATURE

NSF-Simons Center for Quantitative Biology, Northwestern University | June 2021 --- August 2021

- Worked with Drs. Alvin Bayliss and Vladimir Volpert as part of the Quantitative Biology Undergraduate Summer Research Program at Northwestern University.
- Investigated a spatially-explicit (PDE) formulation of the Lotka-Volterra model with two types (competing alleles, species, etc.). We showed that curvature at the interface of regions in which species reside facilitates competitive exclusion and defines the nature, e.g., asymptotic speed, of invasion!

ECOLOGY OF GENETIC INTERVENTIONS AGAINST DISEASE

Departments of Mathematics & Biological Sciences, UI | November 2019 --- May 2022

- With Drs. Chris Remien, Steve Krone, and Jim Bull, investigated the eco-evo-epidemiological consequences of gene drive releases (and other genetic interventions) to suppress vector-borne diseases.
- In work now published in *Evolution, Medicine, and Public Health*, we analyzed mathematical models of human/vector infection dynamics in discrete “patches,” distinguished only by the mode of mosquito biting (i.e., frequency- versus density-dependent mosquito-to-human transmission).

PUBLICATIONS

Yuksel, M. K., Remien, C. H., Karki, B., Bull, J. J., & Krone, S. M. (2021). Vector dynamics influence spatially imperfect genetic interventions against disease. *Evolution, Medicine, and Public Health*, 9(1), 1-10.

Keefer, M., **Yuksel, M. K.**, & Caudill, C. Decision Support Modeling for Outplanting Adult Chinook Salmon in the Santiam River Basin (Technical Report 2022-1 for US Army Corps of Engineers).

TEACHING EXPERIENCE

- **(Co-)instructor** at University of Toronto for the following courses:

Fall 2023		EEB313: Quantitative Methods in R for Biology		32 students
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- **Teaching assistant** at University of Toronto for the following courses:

Winter 2024		EEB313: Quantitative Methods in R for Biology		CD**
Fall 2023		EEB319: Population Ecology		30 students
Fall 2022		EEB458: Evolutionary Quantitative Genetics		8 students
Fall 2022		BIO120: Adaptation & Biodiversity		48 students

- **Teaching assistant** for Dr. Jennifer Johnson-Leung, of the UI Department of Mathematics:

Fall 2021		MATH 461: Abstract Algebra I		17 students
Spring 2021		MATH 215: Proof via Number Theory		16 students
Fall 2020		MATH 215: Proof via Number Theory		15 students
Fall 2019		MATH 330: Linear Algebra		42 students

- Summer 2019, Fall 2021, Spring 2022: **algebra & calculus tutor** at UI Polya Center.

** CD = course development

MENTORSHIP

Summer 2023 Litong Zheng, MSc work-study student
Developed interactive web application of models of viral evolution, spillover, and emergence.

TECHNICAL SKILLS

Advanced: R • \LaTeX
Intermediate: FORTRAN '95 • Python • Julia
Familiar: Mathematica • C

I have particular experience with the following:

- Numerical methods to solve differential equations, e.g., finite-difference and spectral methods.

- Stochastic simulation and computer-intensive statistical methods, including Markov Chain Monte Carlo, expectation-maximization, the Gillespie algorithm, and particle filtering.
- Working with (accessing, normalizing, etc.) genome sequence, methylation, and gene expression data.
- Parallel (multi-core and cluster) computing.
- Tidying, statistical analysis, and visualization of scientific data.
- Shiny app development, including dynamic UI design and reactive programming.

SCHOLARSHIPS, GRANTS, & OTHER AWARDS

John B. George Award (presented to the top graduating senior in the UI College of Science), Spring 2022

College of Science Dean's Award in Mathematics & Statistics, Spring 2022

Outstanding Senior in Department of Mathematics & Statistical Science, Spring 2022

NSF Graduate Research Fellowship Honorable Mention, 2022

College of Science Dean's List, Fall 2018 - Spring 2022

2024-27	CAD\$76,500	University of Toronto Data Sciences Institute Doctoral Fellowship
2023-24	CAD\$10,000	University of Toronto Institute for Pandemics Graduate Studentship
2022-24	CAD\$10,000	University of Toronto Doctoral Recruitment Award
2022-27	CAD\$50,000	University of Toronto Connaught International Doctoral Scholarship
2021-22	USD\$7,594	UI Department of Mathematics Pyrah, Karen & Family Scholarship
2021	USD\$4000	Northwestern Quantitative Biology Undergraduate Summer Research Program
2020-22	USD\$3000	UI College of Science Hill Undergraduate Research Fellowship
2020-21	USD\$3000	UI Department of Mathematics Eugene and Osa Taylor Scholarship
2020	USD\$4161	UI Office of Undergraduate Research Summer Undergraduate Research Fellowship
2019-20	USD\$2222	UI Mathematics Department Scholarship
2019	USD\$1500	UI Department of Biological Sciences Undergraduate Research Grant

PRESENTATIONS

	<u>"A null model for the evolution of the recombination map"</u>
2024	Mathematical Models in Ecology and Evolutionary Biology
	<u>"Spillover: recombination and reservoir life history in viral spillover and emergence"</u>
2024	Evolution, the joint annual conference of the ASN, SSB, and SSE***
2024	UT Annual Microbiology and Infectious Diseases Research Day
2024	UT Institute for Pandemics Annual Symposium
2023	UT EEB Undergraduate Union Graduate Student Series***
2023	Ecology and Evolution of Infectious Diseases (EEID) Conference
2023	UT Atwood Colloquium in Ecology & Evolution***
	<u>"Using patterns of differential methylation to infer relationships between cancer sub-types"</u>
2022	Workshop in Mathematical and Computational Biology*
	<u>"Competitive exclusion by curvature"</u>
2023	UT EEB Undergraduate Union Graduate Student Series***
2022	UI Office of Undergraduate Research Symposium
2022	2022 Pacific Inland Mathematics Undergraduate Conference***

- 2021 UI College of Science Student Research Exposition
 2021 Northwestern Quantitative Biology SURP Symposium***
"Vector dynamics influence spatially imperfect of genetic interventions against disease"
 2021 UI Office of Undergraduate Research Symposium
 2020 UI College of Science Student Research Exposition***
 2020 Idaho Undergraduate Research Conference

*** indicates talk, poster otherwise.

LEADERSHIP & SERVICE

- Ongoing **Ad-hoc reviewer for *Journal of Evolutionary Biology*, G3**
 2024 **University of Toronto Atwood Colloquium in Ecology & Evolution Chair**
 Organized annual Colloquium of students, postdoctoral fellows, and faculty in the Dept. of Ecology & Evolutionary Biology at UofT. Coordinated selection and visits of invited speakers, scheduling of talks and coffee breaks, and a reception which usually concludes the event.
- Fall 2023- **Organizer of University of Toronto EEB Scientific Racism Reading Group**
 Developed and co-lead a reading/discussion group on how research in ecology and evolutionary biology intersects with society (e.g., race and IQ testing, plastic pollution).
- 2023-24 **University of Toronto EEB Student Journal Club and Seminar Co-Chair**
 2023-24 **CUPE 3902 Unit 1 Depy. Ecology & Evolutionary Biology Steward**
 Surveyed and fostered support for Union activities in the Department. Developed a handbook for students in my department, with the goal of clarifying how Union processes (e.g., filing a hiring grievance, filing health insurance claims, etc.) work.
- Winter 2023- **Writer and Associate Editor for The University of Toronto *EEB Quarterly***
 2022-23 **Member of University of Toronto EEB Wellness Committee and EDI Subcommittee**
 Co-led the development of a \$8K undergraduate summer research award for students who self-identify as Black, Indigenous, or People of Colour (BIPOC), and which was not based on GPA. Criteria such as enthusiasm for research and lived experiences used in evaluating applications. In 2023, three students were awarded funds to conduct research in EEB.
- July 2022-Sept 2024 **Volunteer at Toronto Fort York Food Bank**
 2020-21 **Undergraduate member of University of Idaho Coronavirus Committee**
 Identified inconsistencies/gaps in University of Idaho masking, testing, and ventilation policies, and provided COVID policy recommendations to administrators. The work of the committee concluded with the development of a publicly-accessible dashboard with current case counts, hospitalization data, etc. for the University of Idaho and City of Moscow, ID.
- 2017-19 **Member of the City of Moscow, Idaho Sustainable Environment Commission**
 Communicated with the City Council and Mayor about sustainability matters at the University of Idaho, local businesses, etc. The Commission also worked to stop the use of neonicotinoids, a class of pesticides with chemical structure similar to nicotine, in City parks.