

## Tom R. Booker

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CONTACT INFORMATION	University of British Columbia, Vancouver booker@zoology.ubc.ca	778-751-4586 (Cell) <a href="https://tbooker.github.io/">https://tbooker.github.io/</a>
EMPLOYMENT	<p><b>University of British Columbia</b>, Vancouver, Canada</p> <ul style="list-style-type: none"><li>- Bioinformatics Postdoctoral Fellow (Oct 2021 - October 2023)</li></ul> <p><b>University of Calgary</b>, Calgary, Canada</p> <ul style="list-style-type: none"><li>- Postdoctoral Research Fellow (March - October 2021)</li><li>- Supervised by Professor Michael Whitlock and Associate Professor Samuel Yeaman</li></ul> <p><b>University of British Columbia</b>, Vancouver, Canada</p> <ul style="list-style-type: none"><li>- Postdoctoral Research Fellow (Sept 2018 - March 2021)</li><li>- Supervised by Professor Michael Whitlock and Associate Professor Samuel Yeaman</li></ul>	
EDUCATION	<p><b>University of Edinburgh</b>, Edinburgh, Scotland</p> <p>PhD., <a href="#">Evolutionary Genetics</a>, October 2014 - October 2018</p> <ul style="list-style-type: none"><li>- Thesis Title: UNDERSTANDING PATTERNS OF GENETIC DIVERSITY IN THE HOUSE MOUSE GENOME</li><li>- Supervisors: Professor Peter Keightley and Professor Brian Charlesworth</li><li>- I spent Spring 2018 as a visiting student at UBC hosted by Professor Sally Otto</li></ul> <p>MSc., <a href="#">Evolutionary Genetics</a>, 2013 - 2014 (Distinction)</p> <ul style="list-style-type: none"><li>- Thesis Title: SEARCHING FOR BALANCING SELECTION ON A MIMICRY SUPERGENE IN THE BATESIAN MIMIC <i>Papilio polytes</i></li><li>- Supervisors: Professor Deborah Charlesworth and Rob W. Ness (Now Assistant Professor at University of Toronto - Mississauga)</li></ul> <p><b>University of Stirling</b>, Stirling, Scotland</p> <p>BSc Hons, <a href="#">Ecology</a>, 2009 - 2013 (First Class)</p> <ul style="list-style-type: none"><li>- Dissertation Title: AN INVESTIGATION INTO THE FITNESS AND DISTRIBUTION OF A NEWLY DISCOVERED ALLOPOLYPLOID SPECIES, <i>Mimulus peregrinus</i></li><li>- Supervisor: Dr Mario Vallejo-Marin</li><li>- Study abroad at Simon Fraser University, Vancouver, Canada. 2011-2012.</li></ul>	
PUBLICATIONS	<hr/> <p><b>Papers in Revision</b></p> <p><b>Booker, T. R.</b>, Jackson, B., Craig, R., Charlesworth, B. &amp; Keightley, P. D. (<i>In revision at Molecular Biology and Evolution</i>) "Patterns of genetic diversity around protein-coding exons and conserved non-coding elements are explained by strong selective sweeps in mice". Preprint online at: <a href="https://doi.org/10.1101/2021.06.10.447924">https://doi.org/10.1101/2021.06.10.447924</a></p> <p><b>Booker, T. R.</b>, Yeaman, S. &amp; Whitlock, M. C. (<i>In revision at Molecular Ecology Resources</i>) "The WZA: A window-based method for characterizing genotype-environment association". Preprint online at: <a href="https://doi.org/10.1101/2021.06.25.449972">https://doi.org/10.1101/2021.06.25.449972</a></p>	

Grummer, J.\*, Thomaz, A.\* **Booker, T. R.\***, Nietlisbach, P.\*, Matthey-Doret, R.\*, & Whitlock, M. C. (*In revision at Conservation Biology*) “Assisted gene flow is nearly always eventually beneficial, especially when genes are moved in small pulses”.

\* Joint first author

Preprint online at: <https://doi.org/10.1101/2021.04.20.440707>

Exposito-Alonso, M., **Booker, T. R.**, Czech, L., Fukami, T., Gillespie, L., Hateley, S., ... & Zess, E. (*Submitted to bioRxiv*). Quantifying the scale of genetic diversity extinction in the Anthropocene.

Preprint online at: <https://doi.org/10.1101/2021.10.13.464000>

## Published Papers

10. Lind, B. M., Lu, M., Vidakovic, D., Singh, P., **Booker, T. R.**, Yeaman, S., & Aitken, S. (2021 - *In press*). “Haploid, diploid, and pooled exome capture recapitulate features of biology and paralogy in two non-model tree species”. *Molecular Ecology Resources*, 00, 1 14.
9. **Booker, T. R.**, Yeaman S. & Whitlock M. C. (2021). “Global adaptation complicates the search for local adaptation”. *Evolution Letters*, 5: 4-15.
8. **Booker, T. R.**, Yeaman S. & Whitlock M. C. (2020) “Variation in recombination rate affects detection of outliers in genome scans under neutrality”. *Molecular Ecology*, 29: 42744279. *Highlighted on the cover and accompanying perspective piece*
7. Byers K.A., **Booker T. R.**, Combs M., Himsworth C.G., Munshi-South J., Patrick D.M., Whitlock M.C.. (2020) “Using genetic relatedness to understand heterogeneous distributions of urban rat-associated pathogens”. *Evolutionary Applications* 00: 112.
6. **Booker, T. R.** (2020) “Inferring parameters of the distribution of fitness effects of new mutations when beneficial mutations are strongly advantageous and rare”. *G3: Genes, Genomes, Genetics*, 10(7) 2317-2326
5. **Booker, T. R.**, & Keightley, P. D. (2018). “Understanding the factors that shape patterns of nucleotide diversity in the house mouse genome”. *Molecular Biology and Evolution*, 35(12) 2971-2988
4. **Booker, T. R.**, Jackson, B. C., & Keightley, P. D. (2017). “Detecting positive selection in the genome”. *BMC Biology*, 15:98.
3. **Booker, T. R.**, Ness, R. W., & Keightley, P. D. (2017). “The recombination landscape in wild house mice inferred using population genomic data”. *Genetics*, 207(1) 297-309
2. Keightley, P. D., Campos, J. L., **Booker, T. R.**, & Charlesworth, B. (2016). “Inferring the frequency spectrum of derived variants to quantify adaptive molecular evolution in protein-coding genes of *Drosophila melanogaster*”. *Genetics*, 203(2), 975-984.
1. **Booker, T.**, Ness, R. W., & Charlesworth, D. (2015). “Molecular evolution: breakthroughs and mysteries in Batesian mimicry”. *Current Biology*, 25(12), R506-R508.

ACADEMIC HONOURS AND AWARDS	University of British Columbia - Bioinformatics Research Fellowship	2020
	Registration Award - Society of Molecular Biology and Evolution	2019
	<i>Runner up</i> Best student talk at Population Genetics Group 51	2018
	<i>Runner up</i> Best student poster at Population Genetics Group 50	2017
	Environment Yes! <i>Won regional heat - runner up at the national final</i>	Sept 2016
	EASTBIO Doctoral Training Partnership Studentship	2014-2018
	Genetics Society, Sir Kenneth Mather Memorial Prize	2013/2014
	University of Edinburgh, Douglas Falconer Award, best MSc dissertation	2013/2014
	Funding for Undergraduate Summer Project: Botanic Society of Scotland and the Society of Biology	Summer 2012
	<i>Nominated</i> , Simon Fraser University Student Conservation Prize	May 2012
SERVICE & OUTREACH	<b>Reviewing</b>	
	<i>Science, PLoS Genetics, Molecular Biology and Evolution, Evolution, Molecular Ecology, Genome Biology and Evolution, Ecology and Evolution, Frontiers in Zoology, Peer Community in Evolutionary Biology, G3, BMC Biology</i>	
	<b>Academic Service</b>	
	2021 - <i>Present</i> - Member of the Junior Editorial Board at <a href="#">Molecular Ecology</a>	
	2019 & 2021 Undergraduate Student Mentor for Society for Molecular Biology and Evolution	
	<b>Departmental Service</b>	
	2021 - <i>Present</i> - I organise $\Delta$ -tea, an evolutionary genetics discussion group at UBC	
	2020 - <i>Present</i> - Bioinformatics office hours at UBC	
	2020-2021 I ran virtual trivia nights for my department at UBC during the COVID-19 pandemic	
	2020 Postdoc representative on department expansion committee for the Biodiversity Research Centre at UBC	
	2019 - 2020 I co-organised meetings of the Vancouver Evolution Group (VEG) <i>I stopped due to childcare out of work hours</i>	
	2017 - I started and organised a journal club on classic population genetic papers at the University of Edinburgh in 2017	
	<b>Outreach</b>	
	2021 Skype a Scientist participant	
	2020-2021 Essay judge for Canadian Undergraduate Research Competition ( <a href="#">CURC</a> )	
	2020 Mentor for high school students during the Summer of 2020 - organised through Teen Nerd Nite	
	2019 Poster Judge BIOL 310 Animal Behaviour	
	2018-2019 Poster and talk judge EcoEvo Retreat	

TEACHING  
EXPERIENCE

**Course Coordination**

BIOL525D: Bioinformatics for Evolutionary Genetics 2020-2023  
*Upper division course at UBC on bioinformatics.*  
*Website and course materials at: (<https://ubc-biol525d.github.io/>)*

**Teaching Assistance**

Statistics and Data Analysis, MSc course 2014-2017  
*Upper division course, part of the MSc QGGA program at the University of Edinburgh*

Population and Quantitative Genetics, MSc course 2015-2017  
*Upper division course, part of the MSc QGGA program at the University of Edinburgh*

Ecology and Evolutionary Genetics, BSc course 2014-2015  
*Undergraduate course, offered at the University of Edinburgh*

STUDENT  
SUPERVISION

C. Atkinson - Directed studies co-supervisor (2019-2020) - *Undergraduate student at UBC*  
 K.A. Byers - Bioinformatics/genomics mentor for PhD project (2018-2020) - *Now postdoctoral research fellow*  
 S-A. Xerri - Master's project co-supervisor (2018) - *Now PhD student at the Max Planck Institute*  
 C. Barata - Master's project co-supervisor (2017) - *Now PhD student at the University of St. Andrews*  
 B. Lecher - Honour's project co-supervisor (2016-2017) - *Now Pre Doctoral Fellow at the European Bioinformatics Institute*

INVITED  
PRESENTATIONS

**August 2021** - Monash University, School of Biological Sciences Internal Seminar Series, *Online* (Talk)  
*Identifying the genetic basis of local adaptation and testing for convergent evolution across lineages*  
**June 2021** - Virtual SBE 2021 *Online* (Talk)  
*Background selection under evolving recombination rates*  
**May 2021** - Mooers lab group at Simon Fraser University, Canada, *Online* (Talk)  
*The genetics of assisted gene flow*

CONTRIBUTED  
PRESENTATIONS

**March 2021** - EvolTree, *in silico*, Switzerland (Poster)  
*Identifying the genetic basis of local adaptation and testing for convergent evolution across*

*lineages*

**March 2021** - Virtual EcoEvo Retreat, *in silico*, Canada (Talk)  
*Background selection under evolving recombination rates*

**January 2020** - American Society of Naturalists 2020, Asilomar, USA (Talk)  
*Global adaptation confounds the search for local adaptation*

**October 2019** - EcoEvo Retreat, Squamish, Canada (Talk)  
*Leveraging linkage information in studies of local adaptation*

**September 2019** - BLISS, UBC, Vancouver, Canada (Talk)  
*Global adaptation confounds the search for local adaptation*

**July 2019** - SMBE, Manchester, UK (Poster)  
*Patterns of genetic diversity around protein-coding exons and conserved non-coding elements are explained by strong selective sweeps in mice*

**September 2018** - EcoEvo Retreat, Squamish, Canada (Talk)  
*Estimating the parameters of selective sweeps from patterns of diversity around functional elements in wild house mice *Mus musculus castaneus**

**January 2018** - Population Genetics Group 51, Bristol, UK (Talk)  
*Estimating the parameters of selective sweeps from patterns of diversity around functional elements in wild house mice *Mus musculus castaneus**

**August 2017** - ESEB 2017, Groningen, Netherlands (Poster)  
*Selective sweeps and background selection in the genome of wild house mice, *Mus musculus castaneus**

**January 2017** - Population Genetics Group 50, 2017, Cambridge, UK (Poster)  
*Selective sweeps and background selection in the genome of wild house mice, *Mus musculus castaneus**

**July 2016** - SMBE, Gold Coast, Australia (Talk)  
*Hill-Robertson Interference in wild mice, *Mus musculus castaneus**

**December 2015** - Population Genetics Group 49, Edinburgh, UK (Talk - Invited)  
*Hill-Robertson Interference in wild mice, *Mus musculus castaneus**

**July 2015** - SMBE, 2015, Vienna, Austria (Poster)  
*Selective sweeps and background selection in the genome of wild house mice, *Mus musculus castaneus**

**May 2015** - Quantitative Genomics, 2015, London, UK (Talk)  
*Simulating genome evolution in the house mouse: understanding the contribution of Hill-Robertson interference to patterns of genetic diversity*