

CURRICULUM VITAE

William S. Pearman

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EDUCATION

- 2020 – Present PhD Student – Department of Marine Sciences, University of Otago
Advisors: A/Prof Ceridwen Fraser, Dist. Prof. Neil Gemell
- 2017 - 2020 Massey University – School of Natural and Computational Sciences (SNCS), **Master of Natural Science with Distinction**. GPA: 8.3/9
Thesis: Molecular Ecology of an understudied New Zealand Isopod: *Isocladus armatus*
Advisors: Prof. James Dale, Dr. Nicole Freed.
- 2015 - 2017 Massey University, **Bachelor of Natural Science**
Specialty: Molecular Ecology. GPA: 8/9

EMPLOYMENT

- February 2019 - Present **Lab Supervisor**, Massey University, Supervisor: Monika Merriman
Teaching undergraduate labs, coordinating a team of demonstrators.
Course names: Biology of Cells and Biology of Animals. 8-10 hours/week
- February 2018 - Present **Lab Demonstrator**, Massey University, Supervisor: Monika Merriman
Demonstrating lab techniques and teaching undergraduate labs
Course codes: 162.101 and 122.102. 5-6 hours/week
- July 2017 – July 2018 **Research Assistant**, Massey University
Molecular Ecology research assistant, DNA extraction protocol development and performed extractions, bioinformatics pipeline development. 3-6 hours/week

MANUSCRIPTS UNDER REVIEW

- Pearman, W. S.**, Wells, S. J, Silander, O. K., Freed, N. E., & Dales, J (2020). Population structure and dispersal across small and large spatial scales in a direct developing marine isopod *Under review. Equal first author*
- Arranz, V., **Pearman, W. S.**, Aguirre, D. J, & Liggins, L (2020). MARES: a replicable pipeline and curated reference database for marine eukaryote (COI) metabarcoding *Under review. Equal first author*

Pearman, W. S., Freed, N. E., and Silander, O. K., (2020). The advantages and disadvantages of short- and long-read metagenomics to infer bacterial and eukaryotic community composition. *bioRxiv*. <https://doi.org/10.1101/650788>. *Accepted in BMC Bioinformatics*.

Pearman, W. S., Smith, A. N. H., Breckell, G., Dale, J., Freed, N. E., & Silander, O. K. (2018). New tools for diet analyses: nanopore sequencing of metagenomic DNA from stomach contents to quantify diet in an invasive population of rats. *bioRxiv*, 363622. <https://doi.org/10.1101/363622>. *In Review*.

HONORS AND AWARDS

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| 2020 | University of Otago Doctoral Scholarship |
| 2020 | Invited participant: Ira Moana Investigator Workshop
Invitation included full funding for accommodation and registration. The workshop included 13 national and international participants, of which I was one of only 2 students, with all other participants being established academics. Four days spent preparing publications. |
| 2019 | Invited participant: Ira Moana Project Early Career Workshop
Invitation included full funding for accommodation and registration. Three days of data wrangling. |
| 2019 | Best oral presentation: Massey University SNCS Postgraduate Conference. \$100 |
| 2019 | Massey University Masters Scholarship: \$15,000
Awarded based on academic merit and project proposal. |
| 2018 | Presentation Bursary: Oxford Nanopore Conference. \$1000
Awarded to Present at Ecological Society of Australia – 2018 Conference.
Awarded on basis of project merit and novelty. |
| 2018 | Massey Scholarship – Sciences: \$4000
Awarded to top 5% of graduates |
| 2018 | Natural Sciences Scholarship – Masterate Level: \$15,000. Massey University. |
| 2017 | Undergraduate Achievement Award: Massey University
Institute of Natural and Mathematical Science
Awarded to top student in each degree. |
| 2017 | Second Place Prize for Oral Presentation: New Zealand Molecular Ecology Conference |
| 2017 | Conference Travel Award: \$250. New Zealand Molecular Ecology Conference. |

2017 **Second Place Prize in Division for Oral presentations:** Massey University SNCS Postgraduate Conference. \$50.

2015 - 2017 **Natural Sciences Undergraduate Scholarship:** \$15,000. Massey University

CONFERENCE PRESENTATIONS

TALKS

Pearman, W., Freed, N. E., & Silander, O. K., Wells, S. J., Dale, J. (2019). Population genomics of an understudied native marine invertebrate – *Isocladus armatus*. Presented at: *New Zealand Molecular Ecology Conference 2019*. Wellington, New Zealand

Pearman, W., Freed, N. E., & Silander, O. K. (2018). The long and the short of it: Eukaryotic and microbial metagenomics are not the same. Presented at: *New Zealand Molecular Ecology Conference 2018*. Palmerston North, New Zealand

Pearman, W., Smith, A. N. H., Breckell, G., Dale, J., Freed, N. E., & Silander, O. K. (2018). New tools for diet analyses: nanopore sequencing of metagenomic DNA from stomach contents to quantify diet in an invasive population of rats. Presented at: *Ecological Society of Australia 2018*. Brisbane, Australia

Pearman, W., Smith, A. N. H., Breckell, G., Dale, J., Freed, N. E., & Silander, O. K. (2017). New tools for diet analyses: nanopore sequencing of metagenomic DNA from stomach contents to quantify diet in an invasive population of rats. Presented at: *New Zealand Molecular Ecology Conference 2017*. Otago, New Zealand

SKILLS AND EXPERIENCE

1) Bioinformatics

- a. Unix and bash familiarity
- b. *De novo* assembly of genomes
- c. Taxonomic classification with next generation sequencing
- d. Nanopore and Illumina sequence quality processing and demultiplexing
- e. Familiarity with a wide variety of software used for population genomics and metagenomics
 - i. i.e Kraken2, MEGAN, STRUCTURE, minimap2, samtools etc.

2) R Scripting

- a. Data visualization and management, function and pipeline development.

3) Statistics

- a. Population genetic statistics
- 4) Molecular lab experience (i.e PCR, DNA Extraction, protocol development for DNA extraction, etc)
- 5) Library preparation and data processing for next generation sequencing (Nanopore)

NON-ACADEMIC ACTIVITIES

- Scientific outreach to high school students – 2017 – Present
 - Providing molecular lab experience to high school students – 15-20 hours per year
- President of Massey University Albany Community Garden – 2016-2017
- Secretary of Massey Albany Politics Club – 2015-2016
- Volunteer conservation trapper – Okura Bush, 2015-2018
- Recreational hobbies including cooking, 3D printing, and gardening.

CURRENT AND PAST COLLABORATIONS

- Dr. Olin Silander – Comparison of long and short read sequencing for metagenomics
- Dr. Libby Liggins – Metabarcoding database development and analysis pipelines for data re-use in population genetics.

REFEREES

- Dr. Nicole Freed – Academic supervisor, Master degree
 - Senior lecturer in Genetics, Massey University
 - Email: N.Freed@massey.ac.nz
- Professor James Dale – Academic supervisor, Master degree
 - Professor in Evolutionary Ecology, Massey University
 - Email: J.Dale@massey.ac.nz