

# Heath O'Brien, PhD

## Bioinformatician

May 2022

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[Publications](#)

[genome-informatics](#)

## About me

I have 20 years of experience analyzing genetic data, both in academia and the private sector. I've worked on functional genomics, high-throughput sequencing and genotyping data from plants, fungi, bacteria and humans. I enjoy working on interdisciplinary projects and leading teams with diverse skill sets. I also have strong communication skills, having co-authored over 25 publications and presented my research at conferences in 7 countries.

## Skills

Functional genomics

Quantitative genetics

Microbiology

Data visualisation

Scientific communication

Project management

R

Python

BASH

AWS

SQL

## Work History

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|-----------|---|---------------------|
| 2020-2022 | Bioinformatician<br>Theolytics  | Oxford UK           |
|           | <ul style="list-style-type: none"><li>Managing pipelines and compute architecture for genomic analyses of oncolytic viruses</li><li>Advising on research strategy and on the design and analysis of experiments</li><li>Managing partnerships with other companies and academic scientists</li></ul>  |                     |
| 2018-2020 | Lead Bioinformatician<br>Living DNA   | Frome, UK           |
|           | <ul style="list-style-type: none"><li>Leading a team of 6 bioinformaticians and data scientists developing methods to derive ancestry and health insights from genotyping data</li><li>Working to secure external funding to develop novel genomic prediction approaches</li><li>Collaboration with the engineering team to deploy processing pipelines at scale and to improve all steps of sample processing and data delivery</li></ul>                              |                     |
| 2016-2018 | Research Associate<br>MRC Centre for Neuropsychiatric Genetics and Genomics   | Cardiff, UK         |
|           | <ul style="list-style-type: none"><li>Differential gene expression and eQTL analyses on human brain samples</li><li>Integration of functional genomic and large-scale population genetic datasets for schizophrenia and other neuropsychiatric disorders</li><li>Organization of training seminars on data science (Rstudio, MySQL), version control (Git), and pipeline development (Snakemake, Bash)</li><li>Deployment of results on the web using R-Shiny</li></ul> |                     |
| 2013-2016 | Bioinformatician<br>University of Bristol   | Bristol, UK         |
|           | <ul style="list-style-type: none"><li>Comparative transcriptomics to determine the genetic basis of cell wall patterning responsible for colour-production in plant leaves</li><li>Acting group leader during nine-month family leave cover (overseeing a team of three staff and two students, managing procurement)</li></ul>   |                     |
| 2008-2013 | Postdoctoral Fellow<br>University of Toronto  | Toronto, ON, Canada |
|           | <ul style="list-style-type: none"><li>Comparative analyses of bacterial genomes</li><li>Cloning of virulence factors and characterization using in-planta infection assays</li></ul>  |                     |
| 2006-2008 | Lab Administrator<br>Duke University  | Durham, NC, USA     |
|           | <ul style="list-style-type: none"><li>Oversight of Microbiology teaching labs</li><li>Managed one lab technician and eight teaching assistants</li></ul>  |                     |

## Education

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|-----------|--|----------------------|
| 2000-2006 | PhD, Molecular systematics of cyanobacteria<br>Duke University | Durham, NC, USA      |
| 1994-1999 | Bachelors degree, Plant Science<br>University of Alberta       | Edmonton, AB, Canada |

## Training

Advanced Python for Biologists, PR Statistics, February, 2017.  
Advancing in R, PR Statistics, April, 2015.  
Genetic Data Analysis, Summer Institute in Statistical Genetics, June, 2003.

## References

Available on request.