# MICHAEL ROACH

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# **EXPERIENCE**

Jul 2019-Present

Research Scientist, Australian Wine Research Institute

Bioinformatician: grapevine and microbial genomics; interactive data viz; pipeline and webapp dev.

Feb 2016-Jul 2019

Post-Doctoral Research Fellow, Australian Wine Research Institute

Bioinformatician: grapevine and microbial genomics; pipeline dev; protein modelling; molecular docking.

Jul 2015-Feb 2016

Casual, Australian Wine Research Institute

2008-2015

Demonstrator & Tutor, Flinders University

# **EDUCATION**

2015

Doctor of Philosophy, Flinders University

Bioinformatics and biochemistry: enzyme kinetics; protein modelling; molecular docking.

2007

Bachelor of Biotechnology (Hons), Flinders University

# **RESEARCH INTERESTS**

### Genomics

Assembly, comparative genomics, marker discovery, phylogenetics

### Software & Data Viz

- O Bioinformatics tools, pipelines, webapps, interactive data viz
- o bioinf.cc; bitbucket.org/mroachawri; github.com/mroach-awri

### Protein structure

Protein modelling, molecular docking, molecular dynamics

# **SKILLS**

### **Proficient**

Perl, R, Python, Bash, Snakemake, GNU Make, Linux (+ sysadmin), Git (Bitbucket and Github),
 MS Office

### Moderate

o SQL, Fortran, HTML & CSS, Shiny, Django, HPC, Cloud (Azure and Google), Anaconda, Adobe CS

# PRESENTATIONS/AWARDS

### Awards

- O Coding competition winner, BioInfoSummer 2016
- AWRI Recognition Award, 2016

#### **Posters**

- Curating the FALCON-Unzip Chardonnay de novo genome assembly
  - BioInfoSummer 2016
- Population sequencing reveals clonal diversity and ancestral inbreeding in the grapevine cultivar
  Chardonnay
  - 17<sup>th</sup> Australian Wine Industry Technical Conference, 2019
  - Plant and Animal Genome XXVIII Conference, 2020, San Diego, USA

### Talks

- Population sequencing reveals clonal diversity and ancestral inbreeding in the grapevine cultivar Chardonnay,
  - Australian Genomic Technologies Association 2018 Annual Conference
  - 17<sup>th</sup> Australian Wine Industry Technical Conference, 2019; webcast: bit.ly/2ytanb6

# ASSOCIATIONS

### Adelaide Protein Group

- o apg.asn.au
- O Committee member and webmaster

# Australian Society for Biochemistry and Molecular Biology

o asbmb.org.au

### Australian Bioinformatics and Computational Biology Society

o abacbs.org

### **PUBLICATIONS**

Michael J Roach, Daniel L Johnson, Joerg Bohlmann, Hennie JJ van Vuuren, Steven JM Jones, Isak S Pretorius, Simon A Schmidt, Anthony R Borneman (2018) *Population sequencing reveals clonal diversity and ancestral inbreeding in the grapevine cultivar Chardonnay*. PLOS Genetics 14(11): e1007807. https://doi.org/10.1371/journal.pgen.1007807

26 citations\*

Michael J Roach, Simon A Schmidt, Anthony R Borneman (2018) *Purge Haplotigs: allelic contig reassignment for third-gen diploid genome assemblies*. BMC Bioinformatics 19, 460: https://doi.org/10.1186/s12859-018-2485-7 62 citations\*

C Varela, C Bartel, **M Roach**, A Borneman, C Curtin (2019) *Brettanomyces bruxellensis SSU1 Haplotypes Confer Different Levels of Sulfite Tolerance When Expressed in a Saccharomyces cerevisiae SSU1 Null Mutant* Applied and Environmental Microbiology, 85 (4) e02429-18; DOI: 10.1128/AEM.02429-18 4 citations\*

Journal cover! (aem.asm.org/content/85/4.cover-expansion)

Antonio G Cordente, Anthony R Borneman, Caroline Bartel, Dimitra Capone, Mark Solomon, **Michael Roach**, Christopher D Curtin (2019) *Inactivating mutations in Irc7p are common in wine yeasts, attenuating carbon-sulfur 6-lyase activity and volatile sulfur compound production*, Applied and Environmental Microbiology, 85 (6) e02684-18; DOI: 10.1128/AEM.02684-18 2 citations\*

**Michael J Roach**, Anthony R Borneman (2020) *New genome assemblies reveal patterns of domestication and adaptation across Brettanomyces (Dekkera) species*, BMC Genomics 21, 194: https://doi.org/10.1186/s12864-020-6595-z

1 citation\*

**Michael J. Roach**, Anthony R. Borneman and Simon A. Schmidt (2020) *The Origin of Chardonnay clones with historical significance in Australia and California*, Australian Journal of Grape and Wine Research (under review, minor revisions submitted)

# REFEREES

On request

<sup>\*</sup> Google Scholar citation counts as of 18MAY2020