

Xabier Vázquez-Campos, PhD

Computational biologist | Environmental microbiologist

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EDUCATION

2015
|
2011

PhD in Environmental Engineering

The University of New South Wales

📍 Sydney, Australia

• Thesis: "Geomicrobiological aspects of the (bio)leaching of weathered low-grade uranium ore". [doi](#)

2008
|
2007

MSc in Microbiology

Autonomous University of Madrid

📍 Madrid, Spain

• Thesis: "Set-up of molecular ecology methodologies for the monitoring of microbial populations in an indirect reactor for the bioleaching of Zn from polymetallic sulfide concentrates" [in Spanish].

2007
|
2002

Licenciante in Biology

University of Vigo

📍 Vigo, Spain



RESEARCH EXPERIENCE

Current
|
2023

Visiting fellow

School of Biotechnology and Biomolecular Sciences

📍 The University of New South Wales

2022

Senior research associate / Postdoctoral fellow

NSW Systems Biology Initiative. School of Biotechnology and Biomolecular Sciences

📍 The University of New South Wales

2021
|
2017

Research associate / Postdoctoral fellow

NSW Systems Biology Initiative. School of Biotechnology and Biomolecular Sciences

📍 The University of New South Wales

2017
|
2015

Research associate / Postdoctoral fellow

BioGEMS group. Water Research Centre and School of Civil and Environmental Engineering

📍 The University of New South Wales

2011

Research assistant (casual)

BioGEMS group, School of Civil and Environmental Engineering

📍 The University of New South Wales

2011
|
2010

Research assistant

Bioprocesses Group, Faculty of Chemistry

📍 University of Vigo

CONTACT

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🐦 [XabiVC](#)

🌐 [xvazquezc](#)

📺 [xvazquezc](#)

in [LinkedIn](#)

℞ [ResearchGate](#)

🔖 [Google Scholar](#)

🆔 [0000-0003-1134-5058](#)

℞ [E-4772-2013](#)

Σ COMPUTATIONAL BIOLOGY

- Amplicon data analysis (16S, 18S and ITS).
- Genome-based metagenomics.
- Phylogenetics and phylogenomics with maximum likelihood methods.
- Assembly and annotation of (meta)genomes, including eukaryotes.
- Experience with short and long read data.
- Structural modelling of proteins and protein complexes. Protein model evaluation.

> CODING SKILLS

- Main languages: BASH and R. Some SQL and AWK.
- Reproducible reporting with Rmarkdown.
- Environment management with Conda.
- Experienced HPC user.
- Data visualisation and integration.

Made with the *R* package [pagedown](#).

The source code is available on my [datadrivencv-academic repository](#).

Last updated on 2023-03-09.

- 2007
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2006
- **Research assistant**
Bioprocesses Group, Faculty of Chemistry
📍 University of Vigo
 - **Teaching assistant**
Faculty of Biology
📍 University of Vigo



TEACHING EXPERIENCE



Courses

- 2020
- **CVEN9855 - Water & Wastewater Analysis**
School of Civil and Environmental Engineering.
📍 The University of New South Wales



Invited Lectures

- 2012
- **Uranium heap leaching: Microbiological aspects**
School of Civil and Environmental Engineering.
📍 The University of New South Wales




Workshops

- 2019
- **EMBL-ABR: Phylogenetic trees for beginners**
EMBL-ABR (Sydney node).
📍 The University of New South Wales
• Node organiser and facilitator.
 - **EMBL-ABR: Implementing Scalable Bioinformatic Workflows in Snakemake & Nextflow**
EMBL-ABR (Sydney node).
📍 The University of New South Wales
• Facilitator.
- 2018
- **EMBL-ABR: 16S Metagenomics with Galaxy Australia**
EMBL-ABR (Sydney node).
📍 University of Sydney
• Facilitator.
 - **Long-read Data Analysis Workshop**
NSW Systems Biology Initiative.
📍 The University of New South Wales
• Organiser, trainer, facilitator.
- 2011
- **Hands on Zotero, a free alternative to EndNote**
Bioprocesses group.
📍 University of Vigo
• Organiser, sole trainer.

Supervision

Current
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2019


- **Kevin Yonathan**
PhD. Role: External supervisor.  University of Technology, Sydney

Study of the response of microbial communities to silver nanoparticles through metagenomics.

I have also informally supervised, supported, mentored, or collaborated with a total of 21 students across multiple levels (1 Course project, 7 Honours, 12 PhD, and 1 Practicum).


GRANTS

2021
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2020

- **Bioplatforms Australia / NSW Department of Industry RAAP – NCRIS NSW Co-investment Support Shared Grant**
 School of Biotechnology and Biomolecular Sciences / Ramaciotti Centre for Genomics at The University of New South Wales


• Total amount: AU\$600,000

2020

- **BABS sequencing grant**
Investigation of a phage capable of species cross-infection
 School of Biotechnology and Biomolecular Sciences / Ramaciotti Centre for Genomics at The University of New South Wales

• AU\$5,000 for sequencing

2020
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2019

- **ResTech Support Scheme**
OTUreporter pipeline coding project
 The University of New South Wales


• AU\$15,000

2015

- **Postdoctoral writing Fellowship Grant**
 School of Civil and Environmental Engineering at The University of New South Wales


• 0.4 FTE salary for 12 weeks

2015
|
2011

- **UNSW Tuition fee scholarship**
 Graduate Research School at The University of New South Wales

• Tuition fees for PhD program

2015
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2011

- **Australian Postgraduate Award (Industry)**
 Australian Federal Government

• AU\$27,651.00 per year for 3.5 years

AWARDS

2020

- **SRAP Laureate 2020**
 Spanish Researchers in Australia-Pacific



PUBLICATIONS

- 2022 ● **Metaproteomics reveals methyltransferases implicated in dichloromethane and glycine betaine fermentation by ‘*Candidatus Formimonas warabiya*’ strain DCMF**
Frontiers in Microbiology 13, 1035247. [doi](#)
- S. I. Holland, X. Vázquez-Campos, E. Haluk, R. J. Edwards, M. J. Manefield & M. Lee.
- 2021 ● **Community-led, integrated, reproducible multi-omics with anvi'o**
Nature Microbiology 6(1), 3-6. [doi](#)
- A. M. Eren, E. Kiefl, A. Shaiber, I. Veseli, S. E. Miller, M. S. Schechter, I. Fink, J. N. Pan, M. Yousef, E. C. Fogarty, F. Trigodet, A. R. Watson, Ö. C. Esen, R. M. Moore, Q. Clayssen, M. D. Lee, V. Kivenson, E. D. Graham, B. D. Merrill, A. Karkman, D. Blankenberg, J. M. Eppley, A. Sjödin, J. J. Scott, X. Vázquez-Campos, L. J. McKay, E. A. McDaniel, S. L. Stevens, R. E. Anderson, J. Fuessel, A. Fernandez-Guerra, L. Maignien, T. O. Delmont & A. D. Willis.
- 2018 ● **Disentangling the drivers of functional complexity at the metagenomic level in Shark Bay microbial mat microbiomes**
The ISME Journal 12, 2619-2639. [doi](#)
- H. L. Wong, R. A. White III, P. T. Visscher, J. C. Charlesworth, X. Vázquez-Campos & B. P. Burns.
- 2015 ● **Uranium binding mechanisms of the acid-tolerant fungus *Coniochaeta fodinicola***
Environmental Science & Technology 49(14), 8487-8496. [doi](#)
- X. Vázquez-Campos, A. S. Kinsela, R. N. Collins, B. A. Neilan, N. Aoyagi & T. D. Waite.
- 2014 ● ***Fodinomyces uranophilus* gen. nov. sp. nov. and *Coniochaeta fodinicola* sp. nov., two uranium mine inhabiting Ascomycota fungi from northern Australia**
Mycologia 106(6), 1073-1089. [doi](#)
- X. Vázquez-Campos, A. S. Kinsela, T. D. Waite, R. N. Collins & B. A. Neilan.



CONFERENCE CONTRIBUTIONS

- 2019 ● **Unusual DPANN Archaea from a radioactive legacy site**
Australian Microbial Ecology Conference (AusME 2019). [doi](#)
📍 Perth, WA, Australia
- X. Vázquez-Campos, A. S. Kinsela, M. W. Bligh, M. R. Wilkins, T. E. Payne & T. D. Waite [oral presentation].

Selected peer-reviewed publications (5 out of 17). For a full list of my publications, including pre-prints and manuscripts under review, please consult [my long-format CV](#).

Selected abstracts presented at conferences (5 out of 10, including 3 oral presentation and 7 posters).

I have also contributed to 21 abstracts presented by co-authors at conferences.

2018

● **OTUreporter: an automated pipeline for the analysis and report of amplicon sequencing data**

ABACBS-2018 National Conference. [doi](#)

📍 Melbourne, VIC, Australia

X. Vázquez-Campos, A. Chilton, J. Koval, J. J. Tree & M. R. Wilkins [poster].

● **New Archaea lineages from a radioactive legacy site**

Molecular biology of archaea: From mechanisms to ecology (MoBA6). [doi](#)

📍 Vienna, Austria

X. Vázquez-Campos, A. S. Kinsela, M. W. Bligh, M. R. Wilkins, T. E. Payne & T. D. Waite [poster].

2017

● **Metagenomic and geochemical changes following rainfall at a legacy radionuclide waste disposal site**

Australian Microbial Ecology Conference (AusME 2017). [doi](#)

📍 Melbourne, VIC, Australia

X. Vázquez-Campos, A. S. Kinsela, T. E. Payne & T. D. Waite [oral presentation].

2016

● **Rainfall-mediated biogeochemical changes at a legacy radionuclide waste disposal site**

16th International Symposium on Microbial Ecology (ISME16). [doi](#)

📍 Montreal, Canada

X. Vázquez-Campos, A. S. Kinsela, J. J. Harrison, K. Wilsher, S. Thiruvoth, B. Rowling, T. E. Payne & T. D. Waite [poster].



SERVICE

● **Institutional Service**

· **University of New South Wales**

- Postgraduate student representative at the OH&S Committee of the School of Civil and Environmental Engineering (2013–2015).

● Professional Affiliations

- American Society for Microbiology
 - Regular member (2015–Current).
- Australian Bioinformatics and Computational Biology Society
 - Regular member (2018–Current).
- Australian Society for Microbiology
 - Regular member (2015–Current).
- International Society for Microbial Ecology
 - Regular member (2014–Current).
- Spanish Researchers in Australia-Pacific
 - Co-founder (2014).
 - Board member and web/IT coordinator (2014–2020).
 - Regular member (2021–Current).

● Conferences

- Australian Microbial Ecology conference 2022 (AusME2022)
 - Plenary chair and poster judge (2022).
- 6th Australia-Spain Research Forum: Big Data
 - Event registration (2020).
- 5th Australia-Spain Research Forum: 5 years Bridging Australia and Spain in Research and Academia
 - Website (2019).
- 4th Australia-Spain Research Forum: Earth
 - Website (2018).
- 3rd Australia-Spain Research Forum: Air
 - Website and event registration (2017).
- 2nd Australia-Spain Research Forum: Light
 - Website (2016).
- Australia-Spain Water Forum
 - Website and event photography (2015).

● Peer Reviews

Reviewer for >10 different journals since 2016, including leading journals in their own fields, e.g., *Astrobiology*, *Environmental Science and Technology*, or *Microbiome*.

For a full record as reviewer, consult my public [ResearcherID profile](#).