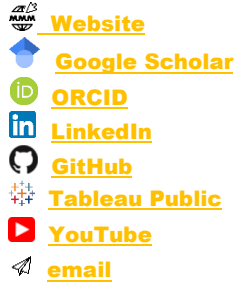


RESUME

Dr. Delphine VINCENT

Data Science, Machine Learning & Bioinformatics

Melbourne area. Dual Citizenship: Australian and French.



SNAPSHOT

- Certifications in **Data Science Machine Learning, Data Analytics, Website Design** and **Chatbot development** (provider: [CodeCademy](#)).
- **PhD** in Plants Science, 4 post-doctoral fellowships in biology, 10 years as a Senior Research Scientist
- 27 years expertise in **biology and data analysis** in academia, and government,
- 2 years as a **data scientist ML / web designer consultant** [ABN 74 267 801 710]
- 62 **publications** (36 peer-reviewed articles, 5 book chapters, 4 editorials, 18 conference papers, 2 patents, 1 preprint, 1 obituary) + 11 **datasets** in public repositories (MassIVE, Pride, (Giga)DB) + 32 GitHub **repositories** + 7 Tableau **vizzes**
- h-index 21, i10-index 29, cited 3,313 times
- in-depth expertise on experimental design, sample collection and preparation, data acquisition, **statistical analyses, machine learning, data mining** and **visualisation**, as well as report **writing**, submission, revision, and publication
- Managerial and **project leading** experiences
- **Editor** and **reviewer** for multiple scientific journals

EMPLOYMENT HISTORY

Senior Research Scientist

- 2013-2023: [Agriculture Victoria Research](#) (AVR), Victorian Government, AgriBio Bundoora VIC 2083, Australia

Post-doctorates

- 2009-2011: [Australian National University](#) (ANU), Canberra ACT 0200, Australia
- 2007-2009: [UMR BIOGECO INRA](#) 69 route d'Arcachon, 33612 Cestas, France
- 2005-2006: [UMR Genetique Vegetale INRA](#), Ferme du Moulon, 91190 Gif-sur-Yvette, France
- 2003-2005: [University of Nevada, Reno](#) (UNR), 1664 North Virginia Street, Reno NV 89557 USA

EDUCATIONAL QUALIFICATION

PhD: 1999-2003: "Proteomic analysis of response to water stress in maize based on cell differentiation". [PhD doctorate](#) granted with Highest Distinctions from [Universite Paris-Saclay, France](#).

DATA SCIENCE SKILLS

Computer OS

Fully skilled with MS Windows software (Word, Excel, Powerpoint, Sharepoint, Teams, OneNote, Outlook, Power BI etc...); familiar with Mac OS systems.

Programming (code languages / IDE)

R/RStudio (tidyverse, readr, tidr, dplyr, ggplot2, Bioconductor), Python/JupyterLab (pandas, NumPy, scikit-learn, SciPy, statsmodels.api, MiniSom, TensorFlow, UpSet, wordcloud, spaCY, Hugging Face Transformers, squarify, RegEx, NLTK, UMAP, gradio, iPython, matplotlib, seaborn, Regex, Pickle, Gensim, pyTorch, XGBoost, LightGBM, Keras, plotpy, BeautifulSoup, Selenium, tqdm, skimage, pillow, imageio, librosa, OpenCV, mido, Pydub, gTTS, MoviePy, Subprocess, FastAPI, Pydantic, RapidFuzz, JSON), Bash Shell, Git, GitHub, Structured Query Language (SQL), Hypertext Markup Language (HTML), javascript (JS), Cascading Style Sheet (CSS), Visual Studio (VS) Code, Anaconda navigator.

Data Science, Machine Learning (ML) and Natural Language Processing (NLP)

Data wrangling, feature engineering, descriptive (means, SDs, quartiles, medians, box plots, histograms, etc..), univariates (linear models, ANOVA, and regression), bivariate (correlation) and multivariates (principal component analyses, partial least square, least discriminant analysis, k-means, self-organising map, and hierarchical clustering analyses) analysis; ML algorithms (Random Forest, Ensemble, Support Vector Machine, Stacked Model, Multi-Layer Perceptron, Deep Learning, Neural Networks), NLP algorithms (tokenisation, stopword removal, stemming, lemmatisation, BoW, TF-IDF, word embedding, POS tagging, topic modelling).

Bioinformatics and Data Mining

BLAST+, Jexpress, Cluster-TreeView, Blast2GO, Goblet, SignalP, TargetP, SecretomeP, ChloroP, NetCGly, GPI-SOM, Goanna PowerBI, Peptide atlas, protein atlas, cytoscape, UniprotKB, String, SwissModel, Apollo Jbrowser, Proteowizard, TransProteomics Pipeline, Pretzel, AgriGO, REVIGO, ShinyGO, Reactome, Pathway Tools, KEGG, Galaxy, BreadwheatCyc, Circos, MetaboAnalyst, DeconTools, EMBL, EBI InterPro, PDBeKB, RCSB PDB Archive, AlphaFold, ColabFold, Cn3D, Bio edit, Mega XI, BLAST, Expasy, Gene Ontology, OpenAI, PaintOmics, Mercator, Galaxy platforms, JBrowser, Integrated Genome Browser.


Data Visualisation

Circos plots, radar plots, violin plots, word clouds, treemaps, treemap bars, histograms, line plots, bar plots, pie charts, scatterplots, donut charts, heat maps, box plots, volcano plots, Gantt chart, LM plots, density plots, join plots, pair plots, pair grid plots, heatmaps, etc.. using PowerBI, Tableau, R and Python.


Data interpretation and Storytelling


Attested by my published articles in which I summarise the process in a simplified schematic diagram, present all results with appropriate charts to facilitate interpretation, as well as by my Jupyter Notebooks and Tableau / PowerBI dashboards.

PUBLICATIONS AND SCRIPTS

Full article list [here](#) and on [Google Scholar](#) .

Python, SQL and HTM/CSS codes on [GitHub](#) .

Tableau dashboards on [Public Tableau](#) .

HTML and CSS code on [website](#) .

DATASETS ON PUBLIC REPOSITORIES

Mass Spectrometry Interactive Virtual Environment (MassIVE):

- 2016-2022: MSV000090572 DOI:10.25345/C53N20J8S, MSV000088253 DOI:10.25345/C5585Q, MSV000085379 DOI:10.25345/C5QQ6J, MSV000084216 DOI:10.25345/C5C95X, MSV000083970 DOI:10.25345/C56S7Z, MSV000083191 DOI:10.25345/C5HG7N, MSV000082070, MSV000081618, MSV000080036

Protein identification database (PRIDE): 2016: PXD002529

(Giga)ⁿ DB: 2023: [Supporting data for "A community resource to mass explore the wheat grain proteome and its application to the Late Maturity Alpha-Amylase \(LMA\) problem"](#)

Apollo Jbrowse: 2024: [Mapping of bread wheat genome, transcriptome and proteome \(proteogenomics\)](#)

CERTIFICATIONS & PROFESSIONAL DEVELOPMENT

CODECADEMY ONLINE COURSE

- 2024: Career Path: Data Scientist – Analytics Specialist (SQL, Python 3, Tableau, Excel, GitHub). [Certificate](#)
- 2024: Skill Path: Build a website with HTML, CSS and GitHub pages. [Certificate](#)
- 2025: Career Path: Data Scientist – Machine Learning Specialist (Python 3). [Certificate](#)
- 2025: Courses: Introduction to Prompt engineering + Introduction to Large Language Model (LLM). [Certificate](#)
- 2025: Skill Path: Build chatbots with Python. [Certificate](#)

TEACHING AND STAFF MANAGEMENT

- 2013-2023: Manager of Science A staff at AVR. Mentoring and supervising students.
- 2013-2020: Demonstrator and lecturer in the Systems Biology Workshop held yearly at AgriBio.
- 2010: Demonstrator in sessions of the Advances in Molecular Plant Science course for undergraduate students at ANU.
- 2007-2008: Guest lecturer in the Biochemistry Master at Bordeaux University (France).

OCCUPATIONAL HEALTH AND SAFETY (OHS), FIRST AID, WELLBEING, MENTORING

- 2013-2016: OHS Representative (HSR), designated first aider (DFA), wellbeing champion for AVR.
- 2022-2023: mentor for the national Young Indigenous Women's STEM Academy (YIWSA) program.

AUSTRALIAN AND INTERNATIONAL COMMITTEES

- 2019-2023: European Cooperation in Science and Technology Action "Protein Production in Photosynthetic Organisms (R3PO)"
- 2021-2023: Proteomics and Metabolomics Victoria (PMV) committee member

EDITORIAL BOARDS

- 2021-2023: BMC Plant Biology editor
- 2015-2023: Editor of the following Research Topics/Special Issues:

[How can secretomics help unravel the secrets of plant-microbe interactions?](#)

[Secretomics: More Secrets to Unravel on Plant-Fungus Interactions. Vol 1](#)

[Secretomics: More Secrets to Unravel on Plant-Fungus Interactions. Vol 2](#)

[Proteomics: Technologies and Their Applications](#)

[Plant Adaptation to Their Biotic and Abiotic Environment Through the Lens of Secretomics](#)

[State-of-the-Art Molecular Plant Sciences in Australia](#)

[Sowing the seed to ensure the future of plant proteomics: Commemorative Issue in Honor of Dr. Dominique Job \(1947-2022\)](#)