

# José Guilherme de Almeida, PhD

Data scientist and ML researcher with 5+ years of experience with clinical data  
*they/them*

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## Work and research positions

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| 2022 → present | <b>Clinical AI Researcher and developer</b><br><i>Champalimaud Foundation</i><br>Lisbon, Portugal <ul style="list-style-type: none"><li>• <b>Team:</b> Computational Clinical Imaging Group (Nickolas Papanikolaou)</li><li>• Developed clinically-valuable deep-learning models for radiological image segmentation and classification</li><li>• Developed self-supervised learning methods to improve classification and segmentation performance using massive datasets</li><li>• Managed and processed large collections of MRI data (&gt;10,000 patients)</li><li>• Developed and implemented clinical image generation methods</li><li>• Collaborated with over 70 researchers across different projects focused on creating value for clinicians and patients</li></ul> |
| 2017 → 2022    | <b>PhD fellow</b><br><i>EMBL-EBI + Cambridge University</i><br>Cambridge, UK <ul style="list-style-type: none"><li>• <b>Advisors:</b> Moritz Gerstung and George S. Vassiliou</li><li>• Developed machine- and deep-learning methods to detect/characterize cells in digitised blood slides and predict disease genetics through uncover cytomorphological profiles</li><li>• Modelled longitudinal sequencing experiments using Bayesian statistics; determined genetic and non-genetic factors driving clonal expansion. Modelled phylogenetic and phylodynamic lifelong trajectories of clones using single-cell colonies</li></ul>   |
| 2016 → 2017    | <b>MSc. Researcher</b><br><i>Center for Neuroscience and Cell Biology</i><br>Coimbra, Portugal <ul style="list-style-type: none"><li>• <b>Advisor:</b> Irina Moreira</li><li>• Developed machine-learning protocols to determine hot-spots (important parts) in the binding interfaces of proteins</li><li>• Performed structural and statistical analysis of large collections of protein-protein complexes and structural characterization of complexes with no known structure</li></ul>  |




















## Education

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| 2017 → 2022 | <b>PhD</b> in computational biology<br><b>University of Cambridge</b> , UK   |
| 2015 → 2017 | <b>MSc</b> in cell and molecular biology (minor in neurosciences)   18/20<br><b>University of Coimbra</b> , Portugal |
| 2012 → 2015 | <b>BSc</b> in biochemistry   16/20<br><b>University of Coimbra</b> , Portugal  |

## Professional training

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| 2024 | Data or Specimens Only Research<br><b>CITI Program (online)</b>                                 |
| 2023 | Data Validation for Machine Learning<br><b>Weights and Biases (online)</b>                      |
| 2023 | Docker & Kubernetes: The Practical Guide<br><b>Academind (online)</b>                           |
| 2022 | Probability theory: foundations for data science<br><b>Colorado Boulder University (online)</b> |
| 2021 | Econometrics: methods and applications<br><b>Erasmus University Rotterdam (online)</b>          |
| 2017 | Summer School in Computational Biology<br><b>University of Coimbra</b>                          |

## Skills, tools and other competencies

Prog. languages	Python 	◆◆◆◆◆
	R 	◆◆◆◆◇
Machine-learning	scikit-learn 	◆◆◆◆◇
	caret 	◆◆◆◆◇
Deep-learning	pytorch  , lightning 	◆◆◆◆◆
	MONAI 	◆◆◆◆◇
	tensorflow 	◆◆◆◆◇
	huggingface 	◆◆◆◆◇
	Large language models (huggingface  , langchain  , ollama)	◆◆◆◆◇
Computer-vision	scikit-image 	◆◆◆◆◇
	OpenCV 	◆◆◆◆◇
Data science	hypothesis testing	◆◆◆◆◇
	multivariate analyses	◆◆◆◆◇
	bayesian methods (MCMC)	◆◆◆◆◇
	Data manipulation (pandas  , polars  , tidyverse  )	◆◆◆◆◇
Data viz	ggplot2 	◆◆◆◆◇
	D3.js (javascript)	◆◆◆◆◇
Web dev	flask 	◆◆◆◆◇
	shiny 	◆◆◆◆◇
	javascript	◆◆◆◆◇
Workflow	version control (git)	◆◆◆◆◇
	containerisation (Docker)	◆◆◆◆◇
	workflow orchestration (snakemake)	◆◆◆◆◇
Soft skills	<p><b>Teamwork:</b> worked with international and pan-European teams on multiple projects</p> <p><b>Leadership, project management:</b> designed and management research projects for students and young researchers</p> <p><b>Communication:</b> clear and precise communication of technical and scientific results to academic and non-specialist audiences</p> <p><b>Adaptability:</b> quickly adapted to new fields, i.e. evolutionary biology and clinical machine-learning</p> <p><b>Critical thinking:</b> identified useful and valuable solutions to complicated problems and implemented them</p>	

## Invited presentations

2024	Bridging imaging and genetics <i>28th Sociedade Portuguesa de Genética Humana Annual Meeting</i> , Porto, Portugal
2024	How does machine-learning assist medical imaging specialists? <i>Data Modelling, AI, and Health: Bridging Models and Insights in Epidemiology, Biostatistics, and Medical Imaging (organized by Mathematics)</i> , Webinar
2023	ProCancer-I: On manufacturer variability, automatic annotation and orphan data <i>European Multidisciplinary Congress on Urological Cancers</i> , Marseille, France

## Conference presentations

2024	Addressing the challenges of curating and segmenting large multi-centric prostate multiparametric MRI datasets with machine-learning (poster) <i>EuSoMII</i> , Vienna, Austria
2024	Predicting Prostate Cancer Biochemical Recurrence After Radical Prostatectomy with Multiparametric MRI Radiomics: A Multicentric Study (poster) <i>EuSoMII</i> , Vienna, Austria

- 2024** Giving new life to orphan data with self-supervised learning - a multi-institutional PCa MRI case study (poster)  
*European Congress of Radiology*, Vienna, Austria
- 2021** The Natural History of Clonal Haematopoiesis (poster)  
*CRUK Cambridge Centre Early Detection Programme 6th Annual Symposium*, Cambridge, UK
- 2020** Leveraging Automated Blood Cell Morphology for Myelodysplastic Syndrome Diagnosis and Prognosis Prediction  
*Quantitative BioImaging Conference*, Oxford, UK
- 2017** Using big-data to understand the protein interface landscape  
*Encontro de Jovens Investigadores de Biologia Computacional Estrutural*, Coimbra, Portugal
- 2016** A Machine Learning Based Protein-Protein Hot-Spot Prediction Method — SpotOn  
*Encontro de Jovens Investigadores de Biologia Computacional Estrutural*, Lisbon, Portugal

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## Honors and Awards

- 2017** EMBL PhD Fellowship (EMBL)
- 2017** MSc honours for outstanding academic performance (Universidade de Coimbra)

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## Teaching experience

- 2024** How to develop a Radiomics Signature  
*ESGAR*, Webinar  
**Description:** Teaching medical professionals about radiomic model development
- 2019** EMBL Lautenschlager Summer School  
*EMBL*, Heidelberg, Germany  
**Description:** Teaching young graduate students about practical bio-image analysis
- 2016** Workshops on Introductory Programming  
*University of Coimbra*, Coimbra, Portugal  
**Description:** Teaching undergraduate students about programming in Python and R

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## Other activities

- 2010 → present** Music (Producer, composer, performer)  
**Description:** Electronic music producer and member of different bands. Member of CIGA239, a musical association in Coimbra.
- 2019** EBI-Sanger-Cambridge PhD Symposium (eSCAMPS) 2019 (Website design)  
**Description:** Developed and designed the website for the 2019 eSCAMPS
- 2018** 20th EMBL PhD Symposium (Organization, speaker contact)  
**Description:** Contacted different high-profile researchers to invite them to present at the 20th EMBL PhD Symposium
- 2014 → 2017** Rádio Universidade de Coimbra (Radioshow host, programming director between 2016 and 2017)  
**Description:** Hosted different radio shows on rock, experimental, jazz and metal music; coordinated the programming department (coordinated a department with approx. 50 people and designed a cohesive radio broadcasting schedule with other departments)
- 2016** Palco RUC (Curator and stage director)  
**Description:** Curated and directed the stage for a small festival attended by thousands of people over four days in Coimbra. Contacted multiple artists and coordinated riders

**2016 → 2017** Junior Enterprise on Science and Technology (Co-founder)  
**Description:** JEST is a junior initiative I founded with a few colleagues that is dedicated to data science training among young students and services to external businesses

## Student orientations

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<b>2024</b>	Retrieval Augmented Generation for Intelligent Medical Chatbot Applications using open-source Large Language Models. <b>João Mata</b> (BSc)
<b>2024 → 2025</b>	Synthetic CT generation for MRI-only prostate radiotherapy. <b>Catarina Caldeira</b> (MSc thesis)
<b>2024 → 2025</b>	Automatic organ-at-risk, clinical target volume and lesion segmentation in prostate radiotherapy. <b>Martim Carneiro</b> (MSc thesis)
<b>2024 → 2025</b>	Any-to-any b-value generation for accelerated diffusion weighted imaging of the prostate. <b>Pedro Parracho</b> (MSc thesis)
<b>2024 → 2025</b>	A systematic study of voxel radiomics and deep-learning for lesion segmentation and detection in prostate cancer bi-parametric MRI. <b>Mariana Ferro</b> (MSc thesis)

## Publications (papers, book chapters, “proceedings”, “other”)

- Almeida, J. G. d., Rodrigues, N. M., Castro Verde, A. S., Mascarenhas Gaivão, A., Bilreiro, C., Santiago, I., Ip, J., Belião, S., Matos, C., Silva, S., Tsiknakis, M., Marias, K., Regge, D., and Papanikolaou, N. (2025), **Impact of Scanner Manufacturer, Endorectal Coil Use, and Clinical Variables on Deep Learning-assisted Prostate Cancer Classification Using Multiparametric MRI**, *Radiology: Artificial Intelligence*, 1, e230555
- Ma, J., Xie, R., Ayyadury, S., Ge, C., Gupta, A., Gupta, R., Gu, S., Zhang, Y., Lee, G., Kim, J., and others (2024), **The multimodality cell segmentation challenge: toward universal solutions**, *Nature Methods*, Nature Publishing Group US New York, 1–11
- Rodrigues, N. M., Almeida, J. G. de, Verde, A. S. C., Gaivão, A. M., Bilreiro, C., Santiago, I., Ip, J., Belião, S., Moreno, R., Matos, C., and others (2024), **Analysis of domain shift in whole prostate gland, zonal and lesions segmentation and detection, using multicentric retrospective data**, *Computers in Biology and Medicine*, Elsevier, 108216
- Almeida, J., Castro Verde, A. S., Gaivão, A., Bilreiro, C., Santiago, I., Ip, J., Belião, S., Matos, C., Tsiknakis, M., Marias, K., and others (2024), **Self-Supervised Learning for Volumetric Imaging: A Prostate Cancer Biparametric Magnetic Resonance Imaging Case Study**, Available at SSRN 4864797
- Rodrigues, N. M., Almeida, J. G. de, Rodrigues, A., Vanneschi, L., Matos, C., Lisitskaya, M. V., Uysal, A., Silva, S., and Papanikolaou, N. (2024), **Deep Learning Features Can Improve Radiomics-Based Prostate Cancer Aggressiveness Prediction**, *JCO Clinical Cancer Informatics*, Wolters Kluwer Health, 8, e2300180
- Del Corso, G., Pachetti, E., Buongiorno, R., Rodrigues, A. C., Germanese, D., Pascali, M. A., Almeida, J., Rodrigues, N., Tsiknakis, M., Papanikolaou, N., and others (2024), “**Radiomics-Based Reliable Predictions of Side Effects After Radiotherapy for Prostate Cancer**”, in *2024 IEEE International Symposium on Biomedical Imaging (ISBI)*, pp. 1–4
- Verde, A. S. C., Almeida, J. G. de, Fonseca, J., Matos, C., Conceição, R. C., and Papanikolaou, N. (2024), “**StitchPro for Computational Pathology Stitching in Patients with Prostate Cancer**”, in *2024 IEEE International Symposium on Biomedical Imaging (ISBI)*, pp. 1–4
- Almeida, J. G. d., Gudgin, E., Besser, M., Dunn, W. G., Cooper, J., Haferlach, T., Vassiliou, G. S., and Gerstung, M. (2023), **Computational analysis of peripheral blood smears detects disease-associated cytomorphologies**, *Nature Communications*, Nature Publishing Group, 14, 4378. <https://doi.org/10.1038/s41467-023-39676-y>
- Rodrigues, A. C., Almeida, J., Rodrigues, N., Moreno, R., Gaivão, A., Bilreiro, C., Santiago, I., Ip, J., Belião, S., Domingues, I., and others (2023), “**Development and Prospective Validation of a Fully Automatic Bi-Parametric MRI Radiomics Signature to Predict Prostate Cancer Disease Aggressiveness: A Multi-Centric Study Using Over 4000 Patients**”
- Almeida, J. G. de, Rodrigues, N. M., Silva, S., and Papanikolaou, N. (2023), **Testing the Segment Anything Model on radiology data**, *arXiv preprint arXiv:2312.12880*
- Rodrigues, N., Almeida, J., and Silva, S. (2023), “**Performance Analysis of Self-Supervised Strategies for Standard Genetic Programming**”, in *Proceedings of the Companion Conference on Genetic and Evolutionary Computation*, pp. 627–630
- Fabre, M. A., Almeida, J. G. d., Fiorillo, E., Mitchell, E., Damaskou, A., Rak, J., Orrù, V., Marongiu, M., Chapman, M. S., Vijayabaskar, M., and others (2022), **The longitudinal dynamics and natural history of clonal haematopoiesis**, *Nature*, Nature Publishing Group, 1–8
- Preto, A. J., Matos-Filipe, P., Almeida, J. G. d., Mourão, J., and Moreira, I. S. (2021), “**Predicting Hot Spots Using a Deep Neural Network Approach**”, *Artificial Neural Networks*, Springer
- Preto, A. J., Barreto, C. A., Baptista, S. J., Almeida, J. G. d., Lemos, A., Melo, A., Cordeiro, M. N. D., Kurkcuoglu, Z., Melo, R., and Moreira, I. S. (2020), **Understanding the binding specificity of G-Protein coupled receptors toward G-proteins and arrestins: Application to the dopamine receptor family**, *Journal of Chemical Information and Modeling*, American Chemical Society, 60, 3969–3984
- R Magalhães, P., Machuqueiro, M., Almeida, J. G. d., Melo, A., DS Cordeiro, M. N., Cabo Verde, S., H Gumus, Z., S Moreira, I., DG Correia, J., and Melo, R. (2019), **Dynamical rearrangement of human epidermal growth factor receptor 2 upon antibody binding: effects on the dimerization**, *Biomolecules*, Multidisciplinary Digital Publishing Institute, 9, 706
- Lemos, A., Melo, R., Preto, A. J., Almeida, J. G., Moreira, I. S., and Dias Soeiro Cordeiro, M. N. (2018), **In silico studies targeting G-protein coupled receptors for drug research against Parkinson's disease**, *Current neuropharmacology*, Bentham Science Publishers, 16, 786–848

- Preto, A. J., Almeida, J. G., Schaarschmidt, J., Xue, L. C., Moreira, I. S., and Bonvin, A. M. (2018), **Computational Tools for the Structural Characterization of Proteins and Their Complexes from Sequence-Evolutionary Data**, *Encyclopedia of Analytical Chemistry: Applications, Theory and Instrumentation*, John Wiley & Sons, Ltd, 1–19
- Melo, R., Lemos, A., Preto, A. J., Almeida, J. G., Correia, J. D., Sensoy, O., and Moreira, I. S. (2018), **Computational approaches in antibody-drug conjugate optimization for targeted cancer therapy**, *Current topics in medicinal chemistry*, Bentham Science Publishers, 18, 1091–1109
- Melo, R., Lemos, A., Preto, A. J., Bueschbell, B., Matos-Filipe, P., Barreto, C., Almeida, J. G., Silva, R. D., Correia, J. D., and Moreira, I. S. (2018), **An Overview of Antiretroviral Agents for Treating HIV Infection in Paediatric Population**, *Current medicinal chemistry*, Bentham Science Publishers
- Moreira, I. S., Koukos, P. I., Melo, R., Almeida, J. G., Preto, A. J., Schaarschmidt, J., Trellet, M., Gumus, Z. H., Costa, J., and Bonvin, A. M. (2017), **SpotOn: high accuracy identification of protein-protein interface hot-spots**, *Scientific reports*, Nature Publishing Group, 7, 1–11
- Almeida, J. G. d., Preto, A. J., Koukos, P. I., Bonvin, A. M., and Moreira, I. S. (2017), **Membrane proteins structures: A review on computational modeling tools**, *Biochimica et Biophysica Acta (BBA)-Biomembranes*, Elsevier, 1859, 2021–2039
- Sensoy, O., Almeida, J. G., Shabbir, J., Moreira, I. S., and Morra, G. (2017), “**Computational studies of G protein-coupled receptor complexes: Structure and dynamics**”, *Methods in Cell Biology*, Academic Press
- Bastos, F. C., Corceiro, V. N., Lopes, S. A., Almeida, J. G. d., Matias, C. M., Dionisio, J. C., Mendes, P. J., Aidos, F. D. Sampaio dos, Quinta-Ferreira, R. M., and Quinta-Ferreira, M. E. (2017), **Effect of tolbutamide on tetraethylammonium-induced postsynaptic zinc signals at hippocampal mossy fiber-CA3 synapses**, *Canadian Journal of Physiology and Pharmacology*, NRC Research Press, 95, 1058–1063
- Melo, R., Almeida, J. G., Verde, S. C., Gumus, Z., Moreira, I., and Correia, J. (2017), “**Structural mechanism of HER2-antibodies complexes by molecular dynamics studies**”, in *Proceedings of MOL2NET 2017, International Conference on Multidisciplinary Sciences, 3rd edition*, p. 5084
- Almeida, J. G., Bonvin, A., and Moreira, I. (2017), “**Using big-data to understand the protein interface landscape**”, in *Proceedings of MOL2NET 2017, International Conference on Multidisciplinary Sciences, 3rd edition*
- Almeida, J. G., Preto, A. J., Melo, R., Gumus, Z. H., Costa, J., Bonvin, A. M., and Moreira, I. S. (2017), “**Coevolution importance on binding Hot-Spot prediction methods**”, in *MOL2NET 2016, International Conference on Multidisciplinary Sciences, 2nd edition*