Date of birth: 15th February 1982

Nationality: French

ORCID: 0000-0001-9006-8436

NIH BioSketch

Github organization: github.com/IARCbioinfo

Website: www.rarecancersgenomics.com

Computational Cancers Genomics Team
Genomic Epidemiology Branch
International Agency for Research on Cancer (IARC-WHO)
25 avenue Tony Garnier, 69007 Lyon, France

Email: follm@iarc.who.int

Research Group: www.iarc.who.int/teams-ccg/

Professional Appointments

2021-on	Team Leader, International Agency for Research on Cancer (IARC-WHO), Lyon, France
2014-2020	Scientist, International Agency for Research on Cancer (IARC-WHO), Lyon, France
2012-2014	Staff Research Scientist, Swiss Federal Institute of Technology, Lausanne, Switzerland
2007-2012	Postdoctoral Researcher, Institute of Ecology and Evolution, University of Berne, Switzerland

Honors and awards

2024-on	Member of the WHO classification of turmours 6 th edition subcommittee on computational pathology, International Agency for Research on Cancer (IARC-WHO), tumourclassification.iarc.who.int/										
2023	Young investigator award, International Mesothelioma Interest Group (iMig), imig.org										
2022-2024	Chair of the Rare Cancers Working Group , European Prospective Investigation into Cancer and Nutrition (EPIC), epic.iarc.who.int										
2022-on	Chair of the Data Science Steering Committee, International Agency for Research on Cancer (IARC-WHO)										
2021-on	Chair of the Health Technologies Pillar , Scientific Steering Committee of the Lyon Auvergne Rhône-Alpes Canceropole (CLARA), www.canceropole-clara.com										
2021-on	lung NET task force member, European NEuroendocrine Tumors Society (ENETS), www.enets.org										
2016-on	Scientific Committee member , French MESOBANK (virtual mesothelioma national biobank), www.netmeso.fr/netmeso/mesopath-et-mesobank										

Education

2004-2007	PhD	in	Population	Genetics	and	Evolution,	University	of	Grenoble,	France.
	theses	ience/tel-00216	6192/							

2001–2004 **MSc Computer science and Applied Mathematics, specialization in Bioinformatics**, Grenoble Institute of Technology ENSIMAG, Grenoble, France.

Personal Statement

I am a computational biologist with extensive training in applied mathematics, computer science, bioinformatics, and statistical genetics. During my doctoral and postdoctoral training, I focused on developing new statistical and computational methods to analyze large genomic datasets, including approaches for Bayesian inference, population genetics, and variant detection. Over the past decade, I have led interdisciplinary teams integrating genomics, transcriptomics, and pathology imaging data to advance translational cancer research. In addition to my role as Team Leader, I also coordinate the bioinformatics efforts at the International Agency for Research on Cancer (IARC-WHO). I am particularly involved in the molecular characterization of thoracic tumors such as lung neuroendocrine neoplasms and malignant pleural mesothelioma; in 2015, I created the Rare Cancers Genomics initiative together with Dr. Lynnette Fernandez-Cuesta. As part of these efforts, I have developed and supervised multiple open-source software projects aimed at ultra-sensitive variant calling, integrative multi-omics pipelines, and, more recently, deep-learning-based image analysis, which have accelerated the discovery of new diagnostic and prognostic biomarkers.

Publications

82 publications in international peer-reviewed scientific journals, including 9 publications as first/co-first author, and 15 publications as last/co-last/corresponding author.

14'406 total citations, h-index: 44 (source: Google Scholar).

Full list available at Google Scholar and My NCBI Bibliography.