Maxime Garcia

PhD in Bioinformatics

8 https://maxulysse.github.io/



Automatization, analysis, reproductibility and visualization

	Experience		
# From 2022	Bioinformatician, Scientific Development Segera Labs	Barcelona, Spain	
	nf-core - Pipeline development;		
= 2016–2022	2022 Bioinformatician , <i>Barntumörbanken Karolinska Institutet & NGI ScilifeLab</i> ♥ Stockholm, Sv		
	nf-core/sarek - Whole Genome/Targeted Sequencing Germline/Somatic analysis	·	
	nf-core/rnafusion - RNA-fusion analysis;		
	nf-core/rnavar - RNA-seq variant analysis;		
= 2013–2014	Research Associate, Cancer Science Institute of Singapore (CSI)	Singapore, Singapore	
	RNAseq analysis; Genome Browser Development.		
= 2009–2013	PhD candidate, Centre de Recherche en Cancérologie de Marseille (CRCM)	■ Marseille, France	
	Development of an Interactome-Transcriptome analysis pipeline; Intern management.		

T Education

	<u>m</u> Education	
= 2013	PhD, Bioinformatics and Genomics, Aix-Marseille Université	₱ Marseille, France
= 2007–2009	Master's Degree, Bioinformatics and Genomics, Université Aix-Marseille II	Marseille, France
= 2007	Erasmus Semester, School of Biological Sciences, University of East Anglia	Norwich, UK
i 2003-2007	Bachelor's Degree, Biology and Biochemistry, Université Aix-Marseille II	Marseille, France

Selected Publications

- [1] JA Fellows Yates and al. "Reproducible, portable, and efficient ancient genome reconstruction with nf-core/eager". In: PeerJ 9.e10947 (2021). DOI: 10.7717/peerj.10947.
- PA Ewels et al. "The nf-core framework for community-curated bioinformatics pipelines". In: Nature Biotechnology 38 (2020), pp. 267-278. DOI: 10.1038/s41587-020-0439-x.
- MU Garcia et al. "Sarek: A portable workflow for whole-genome sequencing analysis of germline and somatic variants [version 2; peer review: 2 approved]". In: F1000Research 9.63 (2020). DOI: 10.12688/f1000research. 16665.2.
- MU Garcia et al. "Interactome-transcriptome integration for predicting distant metastasis in breast cancer". In: Bioinformatics 28.5 (2012), pp. 672-678. DOI: 10.1093/bioinformatics/bts025.

□ Skills

Scripting Groovy, Bash, R Containers Singularity, Docker, Bioconda Web Jekyll, Wordpress

Others Inkscape, LATEX, Gather.town, Markdown

Pipeline Nextflow Management Github, Trello, Slack + HTML5, CSS3, SASS, JS + Linux, Clusters, Git, CI

	Selected Presentations				
= 2020	20ème Journées Ouvertes en Biologie Informatique et Mathématiques (JOBIM)				
= 2019	4th Scandinavian Seminar on Translational Pathology Pergen, Norway				
= 2018	FoUU-day for clinical pathology-cytology				
= 2018	19ème Journées Ouvertes en Biologie Informatique et Mathématiques (JOBIM)				
= 2018	SPHN Workflow Interoperability W	orkshop	Basel, Switzerland		
= 2010	Cancer Bioinformatics Workshop		Cambridge, UK		
	Selected Posters				
= 2018	Keystone Symposia Precision Mede	cine in Cancer	Stockholm, Sweden		
= 2018	Frontiers in Cancer Research and T	herapy 2018	Stockholm, Sweden		
= 2017	European Human Genetics Confere	nce (ESHG) $lacktriangle$ Co	Copenhagen, Denmark		
= 2016	The XVth KI Cancer Retreat		Djurö, Sweden		
= 2013	5th Frontiers in Cancer Science (FC	(S)	Singapore, Singapore		
= 2012	11th European Conference on Computational Biology (ECCB) Basel, Switzerland		Basel, Switzerland		
= 2012	13ème Journées Ouvertes en Biolog	13ème Journées Ouvertes en Biologie Informatique et Mathématiques (JOBIM) Rennes, Fran			
= 2011	12ème Journées Ouvertes en Biolog	rie Informatique et Mathématiques (JOBIM)	Paris, France		
	≡ Organization and Events	5			
# From 2018	nf-core/hackathon	(Member of or	ganization committee)		
= 2015	Journées nationales de la CFD	(Member of organization comm	(Member of organization committee and Webmaster)		
= 2012	Doctoriales en Provence	(Member of or	ganization committee)		
= 2011	Marché de la Chimie - Fête de la S	cience (Conception and	Animation of a stand)		
= 2011	7th BIOTechno forum in Marseille	(Head of or	ganization committee)		
= 2009-2012	Annual meeting of doctoral school	in Biology and Health Sciences from Aix-Mars	eille University		
		(Member of or	ganization committee)		
= 2007-2011	Plan de Cuques French Comics Fes	tival	(Logistic support)		
	Associations				
= 2012-2019	Contributor to 🔞 http://bioinfo	o-fr.net/ (Collaborative French-speakir	ng bioinformatics blog)		
= 2012-2014	Vice-president of JeBiF - RSG-Fran	-president of JeBiF - RSG-France (Association of French Young Bioinformaticians)			
= 2010-2011	President of Hippo'Thèse (Association of young researchers in Biology at Aix-Marseille University)				
= 2010-2011	Vice-president & Webmaster of BIOTechno Network				
	(French Network of Associations organizing forums about careers in Biotechnolo				
= 2009-2010	Webmaster of Hippo'Thèse				
	Languages				
French	Native	English Fluent	Swedish Notions		