Maxime Garcia

PhD in Bioinformatics

☐ 1982-05-14
☐ +46-(0)7-237-343-36
☑ @gau
☐ @MaxUlysse
☑ maxime.garcia@scilifelab.se
② https://maxulysse.github.io/



Automatization, analysis, reproductibility and visualization

	Experience		
🗂 from 2016	rom 2016 Bioinformatician , <i>The Swedish Childhood Tumor Biobank (Barntumörbanken)</i> ♥ Stockholm, Swe Development of a WGS Germline/Somatic analysis workflow in collaboration between SciLifeLab NGI & NE		
	• https://github.com/SciLifeLab/Sarek		
	Whole Genome Sequencing analysis.		
2 013-2014	Research Associate, Cancer Science Institute of Singapore (CSI)	♥ Singapore, Singapore	
	RNAseq analysis; Genome Browser Development.		
2 009–2013	PhD candidate, Centre de Recherche en Cancérologie de Marseille (CRCM)	♥ Marseille, France	
	Development of an Interactome-Transcriptome analysis pipeline; Intern management.		
	<u></u> Education		
2 013	PhD, Bioinformatics and Genomics, Aix-Marseille Université	♀ Marseille, France	
2007-2009	Master's Degree, Bioinformatics and Genomics, Université Aix-Marseille II	♀ Marseille, France	
2 007	Erasmus Semester, School of Biological Sciences, University of East Anglia	♀ Norwich, UK	
2003-2007	Bachelor's Degree, Biology and Biochemistry, Université Aix-Marseille II	♥ Marseille, France	
	₽ Publications		

Publications

- [1] Thomas Bonacci et al. "Identification of New Mechanisms of Cellular Response to Chemotherapy by Tracking Changes in Post-Translational Modifications by Ubiquitin and Ubiquitin-Like Proteins". In: *Journal of Proteome Research* 13.5 (2014), pp. 2478–2494. DOI: 10.1021/pr401258d.
- [2] Maxime Garcia et al. "CNV-Interactome-Transcriptome Integration to detect driver genes in cancerology". In: *Microarray Image and Data Analysis: Theory and Practice.* 2013, pp. 331–338. ISBN: 978-1-46658-682-6.
- [3] Maxime Garcia et al. "Detection of Driver Protein Complexes in Breast Cancer Metastasis by Large-Scale Transcriptome–Interactome Integration". In: *Gene Function Analysis*. 2014, pp. 67–85. DOI: 10.1007/978-1-62703-721-1_5.
- [4] Maxime 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.
- [5] Maxime Garcia et al. "Linking Interactome to Disease: A Network-Based Analysis of Metastatic Relapse in Breast Cancer". In: *Handbook of Research on Computational and Systems Biology: Interdisciplinary Applications*. 2011, pp. 406–427. DOI: 10.4018/978–1–60960–491–2.ch019.
- [6] Tong Liu et al. "Glycosylation Controls Sodium-Calcium Exchanger 3 Sub-Cellular Localization During Cell Cycle". In: European Journal of Cell Biology (2018). DOI: 10.1016/j.ejcb.2018.02.004.

■ □ Skills

Scripting Groovy, Bash, Perl
Containers Singularity, Docker
Web Jekyll, Wordpress
Others Slack, LATEX, Beamer, Inkscape

Pipeline Nextflow, Make Management Github, Trello

+ HTML5, CSS3, SASS, JS

+ Linux, Clusters, Git, CI, R

		♀ Presentations		
	2018	SPHN Workflow Interoperability Workshop	♀ Basel, Switzerland	
	2011	Mathematical and Statistical Aspects of Molecular Biology (MASAMB) XXI	♥ Vienna, Austria	
	2010	Cancer Bioinformatics Workshop	♀ Cambridge, UK	
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		Posters		
	2018	Frontiers in Cancer Research and Therapy 2018	Stockholm, Sweden	
	2017	Department Kick Off (CCK Kick Off)	Stockholm, Sweden	
	2017	European Human Genetics Conference (ESHG)	Copenhagen, Denmark	
	2017	Frontiers in Cancer Research and Therapy 2017	Stockholm, Sweden	
	2016	The XVth KI Cancer Retreat	Q Djurö, Sweden	
	2013	5th Frontiers in Cancer Science (FCS)	Singapore, Singapore	
	2012	11th European Conference on Computational Biology (ECCB)	♀ Basel, Switzerland	
	2012	13ème Journées Ouvertes en Biologie Informatique et Mathématiques (JOBIM)	? Rennes, France	
	2012	6ème Colloque du Cancéropôle PACA		
	2011	12ème Journées Ouvertes en Biologie Informatique et Mathématiques (JOBIM)	P Paris, France	
	2011	19th meeting of doctoral school	♀ Marseille, France	
	2010	Biologistes Chimistes et Physiciens (BCP) VII		
	2010	Interactomics: at the crossroads of biology and bioinformatics	♀ St Raphaël, France	
<u>~~</u>	2009	Fellowhips PhD fellowhip from INSERM/Région PACA		
Ш	2009	The lenowing from INSERM/ Region FACA		
		* Associations		
		Contributor to http://bioinfo-fr.net/ (Collaborative French-speaking bioinformatics blog)		
		Vice-president of JeBiF - RSG-France (Association of French Young Bioinformaticians		
1 2	2010-2011	1 President of Hippo'Thèse (Association of young researchers in Biology at Aix-Marseille University)		
	2010-2011	Vice-president & Webmaster of BIOTechno Network		
~ 2	2009-2010	(French Network of Associations organizing forums about c Webmaster of Hippo'Thèse	areers in Biotechnology)	
		■ Organization and Events		
	2015	Journées nationales de la CFD (Webmaster, Member of	organization committee)	
	2012	Doctoriales en Provence (Member of organization committe	r of organization committee and logistic support)	
	2011	Marché de la Chimie - Fête de la Science (Conception and animation of a stand)		
	2011	7th BIOTechno forum in Marseille (Head of organization committee)		
	2009-2012	Annual meeting of doctoral school in Biology and Health Sciences from Aix-Ma	rseille University	
		(Member of organization committee, abstract selection, jury member, logistic support)		
- 2	007-2011	Plan de Cuques French Comics Festival (Logistic support in organization)		
		□ Languages		
	French	Native English Fluent	Swedish Notions	
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Interests