

# Maxime Garcia

PhD in Bioinformatics

📅 1982-05-14  
☎ +46-(0)7-237-343-36

🐦 @gau

🌐 @MaxUlysse

✉ maxime.garcia@scilifelab.se

🌐 <https://maxulyusse.github.io/>



*Automatization, analysis, reproductibility and visualization*

## 📅 Experience

- 📅 2016-2018 **Bioinformatician**, *Barntumörbanken / Karolinska Institutet* 📍 Stockholm, Sweden  
Development of a WGS Germline/Somatic analysis workflow in collaboration between SciLifeLab NGI & NBIS;  
🌐 <https://github.com/SciLifeLab/Sarek>  
Whole Genome Sequencing 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.

## 🎓 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
- 📅 2003-2007 **Bachelor's Degree**, *Biology and Biochemistry*, Université Aix-Marseille II 📍 Marseille, France









## 📖 Publications

- [1] Tong Liu et al. "Glycosylation Controls Sodium-Calcium Exchanger 3 Sub-Cellular Localization During Cell Cycle". In: *European Journal of Cell Biology* 97.3 (2018), pp. 190–203. DOI: 10.1016/j.ejcb.2018.02.004.
- [2] 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.
- [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. "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.
- [5] 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.
- [6] 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.



















## 💻 Skills

Scripting	Groovy, Bash, Python, Perl, Java, R	Pipeline	Nextflow
Containers	Singularity, Docker, Bioconda	Management	Github, Trello, Slack
Web	Jekyll, Wordpress		+ HTML5, CSS3, SASS, JS
Others	Inkscape, L <sup>A</sup> T <sub>E</sub> X, Beamer, Markdown		+ Linux, Clusters, Git, CI


## Presentations

-  2018 19ème Journées Ouvertes en Biologie Informatique et Mathématiques (JOBIM)  Marseille, France
-  2018 SPHN Workflow Interoperability Workshop  Basel, Switzerland
-  2011 Mathematical and Statistical Aspects of Molecular Biology XXI  Vienna, Austria
-  2010 Cancer Bioinformatics Workshop  Cambridge, UK







## Selected Posters

-  2018 Keystone Symposia Precision Medicine in Cancer  Stockholm, Sweden
-  2018 Frontiers in Cancer Research and Therapy 2018  Stockholm, Sweden
-  2017 European Human Genetics Conference (ESHG)  Copenhagen, Denmark
-  2016 The XVth KI Cancer Retreat  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  Marseille, France
-  2011 12ème Journées Ouvertes en Biologie Informatique et Mathématiques (JOBIM)  Paris, France







## Fellowships

-  2009 PhD fellowship from INSERM/Région PACA

## Associations

-  from 2012 Contributor to  <http://bioinfo-fr.net/> (Collaborative French-speaking bioinformatics blog)
-  2012-2014 Vice-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 Biotechnology)
-  2009-2010 Webmaster of Hippo'Thèse

## Organization and Events

-  2015 Journées nationales de la CFD (Member of organization committee and Webmaster)
-  2012 Doctoriales en Provence (Member of organization committee)
-  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-Marseille University  
(Member of organization committee)
-  2007-2011 Plan de Cuques French Comics Festival (Logistic support)

## Languages

French Native English Fluent Swedish Notions

## Interests

Yoga, Running, Cinema, Poker, Gardening, Ice-Skating, Science-Fiction, Popular science