# **CURRICULUM VITAE**

### **Personal information**

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Keywords

Full Stack Software Engineer, Bioinformatics, Certified Linux System administration, Web applications, Dockers, Linux, Education.



## Professional experience

November 2017 – present

**Manager in Bioinformatics department,** Igenomix, Paterna, Spain.

- ▶ Leading the development for an in-house platform for genomic data visualization and rare disease diagnosis, serving over 50 internal users.
- Certified system administrator for the Bioinformatics HPC infrastructure.
- Developing bioinformatics pipelines for data analysis.

January 2016 - November 2017

Bioinformatics developer at SLU Global Bioinformatics Centre (SGBC), Swedish University of Agricultural Sciences, Uppsala, Sweden.

- Main developer for the *eBioKit 2017 platform* (http://ebiokit.eu/) a stand-alone educational platform for bioinformatics research and training in a controlled environment.
- Main developer in the bioinformatics work package for the B3Africa project (http://www.b3africa.org). This work package includes several tasks such as the implementation of userfriendly and web-based tools for automatized execution of bioinformatics pipelines and the compilation of available bioinformatics tools and databases for omics data analysis.
- ▶ Main developer for *Galaksio*, a user-friendly and workflow-based wrapper for Galaxy for education purposes.

October 2012 - December 2015

**Bioinformatician at the Laboratory of Genomics of Gene Expression**, Príncipe Felipe Research Center, Valencia, Spain.

- ▶ Main developer for *PaintOmics 3*, an integrative web-based visualization platform for multi-omics experiments.
- Main developer for STATegra EMS, an integrative system for multi-omics experiments annotation and tracking.

Bioinformatics Programmer at SLU Global Bioinformatics Centre (SGBC), Swedish University of Agricultural Sciences, Uppsala, Sweden.

## **Education**

- 2024: Linux Foundation training: Kubernetes Fundamentals (LFS258) [Enrolled]
  Linux Foundation training: Introduction to Kubernetes (FS158x).
- ▶ 2022: Linux Foundation Certified System Administrator (LFCS). The Linux Foundation.
- ▶ 2013-2017: PhD at the Universitat Politècnica de València, Valencia, Spain.
  - ▶ Thesis title: Development of bioinformatics resources for the integrative analysis of Next Generation omics data
  - ▶ Thesis supervisors: Dr. Ana Conesa Cegarra, Genomics of Gene Expression Lab, Príncipe Felipe Research Center, and Dr. Monserrat Robles Viejo, Universitat Politècnica de València.
- 2011-2012: Master in Bioinformatics and Computational Biology at the Universidad Complutense de Madrid. Spain.
  - Thesis title: Functional characterization of long non-coding RNA using NGS data.
  - ▶ Project supervisor: Dr. Ana Conesa, Genomics of Gene Expression Lab, Príncipe Felipe Research Center, Valencia, Spain.
- ▶ 2005-2011: Computer engineering at the Universidad Complutense de Madrid. Spain.

## Technical skills

- Computing skills
  - Advanced scripting languages (Python, Bash).
  - ▶ Containerization technologies (Docker, Apptainer [former Singularity]).
  - ▶ UNIX-based systems management (CentOS, Ubuntu, Ansible, Bash).
  - Integrative data visualization techniques.
  - Design and management of AWS architectures.
  - ▶ Web Programming (JavaScript, HTML5, CSS3, Python, Java Servlets, PHP).
  - Design and management of databases (MongoDB, MySQL, PostgreSQL).
  - Deploy and management of CMS (Wordpress, Moodle)
  - Object Oriented Programming (Python, Java).
  - Project management, software testing and documentation.
- Other skills
  - Experience in MS Office tools.
  - Skills in graphic design.
  - Innovative and creative mindset.
  - Initiative-driven with strong organizational skills.
  - Great interpersonal skills, fostering positive relationships within the workplace.

## **Relevant merits**

- Publications
  - Hernández-de-Diego R, Tarazona S, Martínez-Mira C, Balzano-Nogueira L, Furió-Tarí P, Pappas GJ Jr, Conesa A. PaintOmics 3: a web resource for the pathway analysis and visualization of multi-omics data. Nucleic Acids Res. 2018 Jul 2;46(W1):W503-W509.
  - ▶ Hernández-de-Diego R, de Villiers EP, Klingström T, Gourlé H, Conesa A, Bongcam-Rudloff E. The eBioKit, a stand-alone educational platform for bioinformatics. PLoS Comput Biol. 2017 Sep 14;13(9):e1005616.
  - Klingström T, Hernández-de-Diego R, and Bongcam-Rudloff E. Galaksio, a user friendly workflow-

- centric front end for Galaxy. EMBnet.journal, [S.I.], v. 23, p. e897, nov. 2017. ISSN 2226-6089.
- ▶ Hernández-de-Diego R, Boix-Chova N, Gómez-Cabrero D, Tegner J, Abugessaisa I, Conesa A. STATegra EMS: an Experiment Management System for complex next-generation omics experiments. BMC Systems Biology 2014, 8(Suppl 2):S9
- ▶ Conesa A and Hernández-de-Diego R. *Omics Data Integration in Systems Biology: Methods and Applications*. Comprehensive Analytical Chemistry 2014 (64). *doi:10.1016/B978-0-444-62650-9.00016-6*

#### Grants

- ▶ 2016 IRSES- DEANN Marie Curie EU-Latin American for Genomics Research [Grant]
  - December 2016: Guest researcher at the Department of Bioinformatics at the Instituto Nacional de Tecnología Agropecuaria (INTA), USA.
- ▶ 2015 IRSES- DEANN Marie Curie EU-Latin American for Genomics Research [Grant]
  - November December 2015: Guest researcher at the Department Cellular Microbiology, University of Florida, USA.
- ▶ 2014 IRSES- DEANN Marie Curie EU-Latin American for Genomics Research [Grant]
  - May August 2014: Guest researcher at the Department of Computational Genomics, University of Brasilia, Brazil.

#### International conferences

- PaintOmics 3.0: Integrated visualization of multi omics data on KEGG pathways. SMODIA 2015, Valencia, Spain. Oral presentation. September 2015.
- ► The STATegraEMS, an Experiment Management System for multi-omics experiments. SMODIA 2014, Heraklion, Crete, Greece. Oral presentation. November 2014.
- ▶ STATegra EMS: an Experiment Management System for complex next-generation omics experiments. XII Symposium on Bioinformatics, Sevilla. Poster. September 2014
- ▶ The STATegra NGS Experiment Management System. HiTSEQ ISCB/ECCB 2013, Berlin, Germany. Poster. July 2013.
- Developing a NGS Experiment Management System, The STATegra EMS. The Next NGS Challenge Conference, Valencia, Spain. Poster. May 2013.

#### Courses and seminars (teaching)

- ▶ The Genomics of Gene Expression RNA-Seq Course. CIPF, Valencia, Spain. March 2015.
- ▶ The H3ABioNet Advanced Systems Administration workshop: Introduction to the eBioKit system: services, architecture and administration. Pretoria University, Pretoria, South Africa. February 2015.
- The Genomics of Gene Expression RNA-Seg Course. CIPF, Valencia, Spain. September 2014.
- International Course on Massive Data Analysis: Transcriptomics. CIPF, Valencia, Spain, March 2014.
- Introduction to Bioinformatics using the eBioKit platform. KEMRI Wellcome Trust Institute, Kilifi, Kenya. Teacher assistant. August 2013.
- ▶ H3ABioNet Introduction to Bioinformatics using the eBioKit platform. International Center for Insect Physiology and Ecology (ICIPE), Nairobi, Kenya. Teacher assistant. July 2013.
- ▶ EMBNET eBioKit Introduction to the new eBioKit functionalities. ISCB Africa ASBCB Conference on Bioinformatics 2013. Casablanca, Morocco. Teacher assistant. March 2013.