David Mas-Ponte

Bioinformatician

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Education

2015–2017 M.Sc. in Bioinformatics for Health Science

Universitat Pompeu Fabra, Barcelona, Spain.

Specialization in Genomics - GPA: 9 / 10

2014 Exchange Student

McGill University, Montreal, Canada.

Academic international exchange

2011–2015 B.Sc. in Biotechnology

Universitat Autonoma de Barcelona, Spain.

GPA: 8.5 / 10

Research Experience

Jul 2016-today | Centre for Genomic Regulation (CRG) - Master Science Research Thesis

My Master Thesis is focused in the link between IncRNAs' subcellular localization and their function. I have also developed a web-based DB using R and SQL to make subcellular expression data available to the scientific community. **Roderic Guigo's Lab,**

tutored by Rory Johnson.

2015-2016 Institute of Evolutionary Biology (IBE) - Part time Research Internship

I studied the evolutionary processes surrounding the RHD gene in Western Mediterranean populations in order to unravel demographic (*drift*) or adaptive (*selection*) pro-

cesses round this locus. David Comas' Lab

06-08 2014 | Molecular Biology Institute of Barcelona (IBMB) - Research Internship

I took part in the study of the CIC (CaplCua) protein in Drosophila development. I gained research skills in fruit y genetics, in in situ hybridization and in recombinant DNA tech-

niques for CRISPR/Cas9 set up. Gerardo Jimenez's Lab

06-07 2012 Centre for Research in Agricultural Genomics (CRAG)- Research Internship

I participated in the study of casual mutations from a group of neuronal intermediate lament inclusion disease (NIFID) patients and its phenotypic significance. I also took part in a sequencing of a region in transgenic mice which developed a similar phenotype.

Tutored by Alex Clop in the Animal Genomic Department.

Code

Good Python, SQL, R & git
Fair Perl, Shell & LATEX
Learning Ruby & JavaScript (D3)

Publications & Conferences

2017

David Mas-Ponte, Joana Carlevaro-Fita, Emilio Palumbo, Toni Hermoso, Roderic Guigo, and Rory Johnson. Lncatlas database for subcellular localisation of long noncoding rnas. 2017 - Submitted

2016

Andre Flores-Bello, David Mas, Miruna Rosu, and David Comas. Genetic analysis and evolution of rh blood group system in basques. In 6th international conference of the series: DNA Polymorphisms in Human Populations, Musée de l'Homme, Paris, December 2016 - Peer-reviewed Conference - Poster

Languages

Catalan Mother tongue
Spanish Bilingual Proficiency
English Fluent (FCE 2011)

Scholarships & Awards

2015-2016 | MECD - COLAB - Fellowship

I was awarded with the Beca de Colaboracion (Collaboration Fellowship) in Department of Experimental and Health Sciences UPF by Ministerio de Educacion, Cultura y Deporte to collaborate in David Comas Lab.

Sep-Oct 2015 JAE INTRO - CSIC - Fellowship

I was awarded with a JAE INTRO fellowship in Institute of Evolutionary Biology by Consejo Superior de Investigaciones Científicas (CSIC) to collaborate in David Comas Lab.

2014-2015 UAB + AGAUR Exchange - Scholarship

I was awarded with a scholarship from my university and from the Catalan government for my Exchange Academic Experience in CANADA.

Jul-Ago 2014 Passa lestiu al parc-PCB - Scholarship

I was awarded with the scholarship Passa lestiu al parc by Barcelona Science Park (PCB) to collaborate in Gerardo Jimenezs Lab.

2013 VI Premi de Recerca en Ciencies de la Salut-PRBB - Award

I received the VI Premi de Recerca en Ciencies de la Salut (Research Award in Health Science) by Research Park of Biomedicine in Barcelona (PRBB) for my high-school final research project "*Polymorphism in the CCR5 gene*".