# Sergio Castillo Lara

## Bioinformatician

#### **CONTACT**

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Date of birth October 27th, 1992

Nationality Spanish

## **EXPERIENCE**

#### 2017 - Present

## Ph.D. Fellow

Computational Genomics Lab, University of Barcelona

- Development of web-based tools to integrate protein-protein interactions, sequence annotations, and gene expression data for the planaria *Schmidtea mediterranea*.
- Analysis of RNA-seq data for *Schmidtea mediterranea*.
- Development of tools that utilize machine-learning for predicting and retrieving protein-protein interactions.
- Assistant teacher of the courses:

"Introduction to Algorithmics": Msc. Bioinformatics for Health Sciences at UPF.

"Bioinformatics": BSc. Biochemistry at UB.

"Computational Genomics": BSc. Bioinformatics at UPF/UPC/UB.

## 2014 - 2017

## Research Intern

Computational Genomics Lab, University of Barcelona

- Prediction of an interactions network of planarian genes and proteins using sequence homology information and machine learning.
- –Implementation of a pipeline to retrieve protein-protein interactions from scientific literature using natural language processing.

## **EDUCATION**

2015 - 2017

**MSc Bioinformatics for health sciences** 

Universitat Pompeu Fabra, Barcelona, Spain

2010 - 2015

**BSc Biology** 

Universitat de Barcelona, Barcelona, Spain

## PROFESSIONAL SKILLS

Programming Languages Python, Perl, R, JavaScript, Bash

Databases Neo4j, Cypher Query Language, SQL

Data Science ggplot2, scikit-learn, R caret

Web HTML5, CSS, Django, jQuery, Bootstrap, Cytoscape.js

Other Statistics, Machine-Learning, RNA-seq, Linux, git, LaTeX

S. Castillo-Lara, J.F. Abril (2018)

PPaxe: easy extraction of protein occurrence and interactions from the scientific literature. *Bioinformatics*, AOP. doi:10.1093/bioinformatics/bty988

S. Castillo-Lara., J.F. Abril (2018)

PlanNET: Homology-based predicted interactome for multiple planarian transcriptomes. *Bioinformatics*, 34(6): pp. 1016-1023. doi:10.1093/bioinformatics/btx738

D. Boloc, **S. Castillo-Lara**, G. Marfany, R. Gonzàlez-Duarte, J.F. Abril (2015)

Distilling a visual network of retinitis pigmentosa gene-protein interactions to uncover new disease candidates. *PLOS ONE*, 10(8): e0135307. doi:10.1371/journal.pone.0135307

#### POSTERS AND COMMUNICATIONS

S. Castillo-Lara, J.F. Abril (2018)

PPaxe facilitates human-curation of protein-protein interactions filtered from the scientific literature. VI Jornada de Bioinformàtica i Genòmica, Barcelona.

- R. Arenas-Galnares, **S. Castillo-Lara**, V. Toulis, G. Marfany, R. Gonzàlez-Duarte, J.F. Abril (2018) **RPGeNet v2.0 Enhanced navigation through the retinitis pigmentosa interaction network** VI Jornada de Bioinformàtica i Genòmica, Barcelona.
- E. Pascual-Carreras, M. Marín-Barba, **S. Castillo-Lara**, M. Sureda-Gómez, G. Rodríguez-Esteban, H. Heyn, J.F. Abril, E. Saló, T. Adell (2017)

Understanding the anterior and the posterior signaling centers in planarians. XXIX Developmental Biology Meeting, Barcelona.

S. Castillo-Lara, J.F. Abril (2017)

A human-planarian interologs network to annotate planarian transcriptomes: PlanNET 15th International Workshop RECOMB Comparative Genomics, Barcelona.

S. Castillo-Lara, D. Boloc, G. Marfany, R. Gonzàlez-Duarte, J.F. Abril (2015)

Distilling a network of interactions to uncover genes involved in Retinitis Pigmentosa disease. Il Jornada de Bioinformàtica i Biologia Computacional, Barcelona.

## **SCHOLARSHIPS**

| 2017 | Grant for the recruitment of early-stage research staff (FI)<br>Agència de Gestió d'Ajuts Universitaris i de Recerca |
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| 2016 | Beca de colaboración con departamentos (Collaboration Scholarship)<br>Ministerio de Educación Cultura y Deporte      |
| 2014 | Summer Research in Genetics University of Barcelona  |