

ADRIÀ MITJAVILA VENTURA

Bioinformatician at *Istituto Europeo di Oncologia*

Biotechnologist living around computers. Interested in data analysis to support experimental research as well as in pure computational experiments and in software/pipeline development.

References available upon request.



EXPERIENCE

07/2019
|
Present

Bioinformatician

Istituto Europeo di Oncologia

📍 Milan, Italy

Pasini's lab

Analysis of omics data analysis (RNA-seq, ChIP-seq, ATAC-seq) | Comparison of different ATAC-seq peak callers. | Development of an ATAC-seq pipeline for Snakemake.

07/2018
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10/2018

Research Assistant

Instituto de Biomedicina de Sevilla

📍 Seville, Spain

Gene expression lab

Analysis of MNase-seq data from Ccr4-mutant yeast.

09/2017
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07/2018

Master's thesis student

Instituto de Biomedicina de Sevilla

📍 Seville, Spain

Gene expression lab

Analysis of MNase-seq data from Xrn1-mutant yeast | MNase-qPCR experiments, including MNase DNA digestion, primer desing, qPCRs and analysis.

- Supervisors: Dr. Sebastián Chávez de Diego and Dr. Gabriel Gutierrez Pozo.
- MSc degree in Molecular Genetics and Biotechnology.

07/2017
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09/2017

Internship student

Institut d'Investigació Biomèdica de Girona

📍 Girona, Spain

Unit of Diabetes, Endocrinology and Nutrition

Identification of new factors (genetic and bacterial) related to adipose tissue physiology and obesity.

01/2017
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06/2017

Bachelor's thesis student

Microbial. Sistemes i aplicacions analítiques, S.L.

📍 Girona, Spain

DNA extraction of intestinal mucosa biopsies | qPCR experiments to detect clusters and families of bacteria | NGS data analysis to quantify bacterial genders in intestinal biopsies.

- Supervisor: Prof. Jesús García-Gil.
- BSc degree in Biotechnology.

09/2016
|
06/2017

Internship student

Microbial. Sistemes i aplicacions analítiques, S.L.

📍 Girona, Spain

Microbiological and molecular analyses of food and water samples to search for pathogens and perform food forensics.

CONTACT

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✉ adria.mitjavila94@gmail.com
🌐 amitjavilaventura.github.io
🔗 [/amitjavilaventura](https://github.com/amitjavilaventura)
📞 0000-0002-0337-1929
🔗 BioBit

SKILLS

Omics data analysis:

Microarray
RNA-seq
ChIP-seq
ATAC-seq
MNase-seq

Programming/scripting:

R/Bioconductor
Bash
Python

Other:

Snakemake
Conda
Docker
Git

Experimental:

DNA extraction...
(q)PCR
Electrophoresis
Western blot
MNase-digestion
Bacterial/yeast culture

Languages:

Catalan (Native)
Spanish (Native)
English (IELTS 7.5 - C1)
Italian (Basic)



EDUCATION

2018
|
Present

MSc in Bioinformatics and Biostatistics

Universitat Oberta de Catalunya

📍 Online

2017
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2018

MSc in Molecular Genetics and Biotechnology

Universidad de Sevilla

📍 Seville, Spain

- Specialization on Molecular Biology R&D
- Master's thesis: *Influence of Xrn1 in nucleosome positioning across Saccharomyces cerevisiae genome and its effects in transcription.*

2012
|
2017

BSc in Biotechnology

Universitat de Girona

📍 Girona, Spain

- Specialization in Cellular & Molecular Biotechnology.
- Bachelor's thesis: *Butyrate-induced changes in the diversity of intestinal mucosa-associated microbiota in colorectal cancer patients submitted to a lateral ileostomy.*



PUBLICATIONS

2020

Xrn1 influence on gene transcription results from the combination of general effects on elongating RNA pol II and gene-specific chromatin configuration

Begley V *et al.* RNA Biology. DOI: [10.1080/15476286.2020.1845504](https://doi.org/10.1080/15476286.2020.1845504).

DEVELOPMENT

chromHMMviewR:

Visualize outputs from ChromHMM in R.

DPomics (*ongoing*):

Shiny app for omics visualization.

DNAme:

Shiny app to convert your name to a DNA sequence.

ATAC-seq snakemake (*ongoing*):

Snakemake pipeline for ATAC-seq data.

ORGANIZATIONS

ASBTEC:

Board member (2019 - Present)
 Collaborator (2018 - Present)
 Member (2017 - Present)

SCB:

Member (2018 - Present)

HONORS & AWARDS

Elite sportsman:

Federación Española de Remo
2015 - 2021

World Rowing U23 Champion:

BLM2x
Rotterdam, Netherlands.
2016