ADRIÀ MITJAVILA VENTURA

Bioinformatician at Istituto Europeo di Oncologia | MSc

Biotechnologist living around computers, interested in data analysis to support experimental research as well as in pure (bio)computational experiments and in software/pipeline development.



EXPERIENCE

07/2019 | Present

Bioinformatician

Istituto Europeo di Oncologia

Milan, Italy

- \bullet Analysis and integration of RNA-seq, ChIP-seq and ATAC-seq data.
- Development of an ATAC-seq pipeline for Snakemake.

09/2017 | 10/2018

Master's thesis student & Research assistant

Instituto de Biomedicina de Sevilla

Seville, Spain

- · Analysis of MNase-seq data.
- Performance and analysis of MNase-qPCR experiments.

09/2016 | 06/2017

Internship & Bachelor's thesis student

Microbial. Sistemes i aplicacions analítiques, S.L.

Girona, Spain

- qPCR of intestinal biopsies to detect/quantify bacterial genres.
- Statistical analysis of high-throughput sequencing data to quantify bacteria species.



EDUCATION

2018 | Present MSc in Bioinformatics and Biostatistics

Universitat Oberta de Catalunya

Online

2017 | 2018

MSc in Molecular Genetics and Biotechnology

Universidad de Sevilla

Seville, Spain

 Master's thesis: Influence of Xrn1 in nucleosome positioning across <u>Saccharomyces</u> <u>cerevisiae</u> genome and its effects in transcription.

2012 | 2017

BSc in Biotechnology

Universitat de Girona

Girona, Spain

• Bachelor's thesis: Butyrate-induced changes in the diversity of intestinal mucosaassociated 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.



CONTACT

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- amitjavilaventura.github.io
- (7) /amitjavilaventura

SKILLS

Omics data analysis:

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

Programming/scripting:

R/Bioconductor Python Bash

Other:

Snakemake Conda Docker Git

Languages:

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

> Build with pagedown Source code available here Last updated on 2021-04-13



• Rotterdam, Netherlands.